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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Title:
Perfect score:
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Gapop 10.0 , Gapext 0.5
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482.536 Million cell updates/sec
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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             US-08-484-408A-44
US-08-105-483-214
US-08-709-209-214
US-08-105-483-216
US-08-105-483-216
US-08-105-483-216
US-08-709-209-216
US-08-709-209-216
US-08-458-101-219
US-08-458-101-219
US-08-486-099-106
US-08-486-099-116
US-08-486-107-116
US-08-486-107-106
US-08-486-2551A-106
US-08-485-551A-106
US-08-471-913A-106
US-08-481-264A-106
US-08-361-707-95
US-09-361-707-95
US-08-480-173A-45
US-08-480-173A-45
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     Sequence 106,
Sequence 93, 2
Sequence 95, 2
Sequence 95, 2
Sequence 95, 2
Sequence 95, 2
Sequence 45, 2
Sequence 45, 2
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Sequence
Sequence
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     97, Appl
44, Appl
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214, Appl
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216, App
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219, App
210, App
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Matches 12
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US-09-361-707-97
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Patent No. 6258937
GENERAL INFORMATION:
APPLICANT: Tong,
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                                          1 PLSSIFSRIGDP 12
     PLSSIFSRIGDP
                                                                                                  1 Similarity
12; Conserv
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                                                                                               Conservative
52
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61-707-4	-361-707-4	-08-484-408A-	US-08-480-173A-48	-08-683-262B-	US-08-683-262B-48	-08-683-	-09-361		707-9	US-09-361-707-89	US-09-361-707-87	- 1	-09-361-707-		•	Ñ	5204096-1	ľ	
48	e 46	48	48	Sequence 56, Appl	40	D (4	49	90	00	e 87.	85,	•	Facent No. 5164485	10. 020109	5	•	Sequence 55, Appl	

ALIGNMENTS

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; SEQUENCE DESCRIPTION: SEQ ID NO: 97: US-09-361-707-97
                                                                                                                                                                                        TELEPHONE: (617) 542-5070
TELEPHAX: (617) 542-8906
INFORMATION FOR SEQ ID NO: 97:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                 SOFTWARE: FASESEQ for Windows Version 2.
CURRENT APPLICATION NUMBER: US/09/361,707
APPLICATION NUMBER: US/09/361,707
FILING DATE: 27-Jul-1999
PRIOR APPLICATION NUMBER: 08/683,262
APPLICATION NUMBER: 08/683,262
FILING DATE: 18-JUL-1996
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 34,310
REFERENCE/DOCKET NUMBER: 00786/287003
TELECOMMUNICATION INFORMATION:
TRIREPHONE: 1617,625-676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: MA
COUNTRY: USA
ZIP: 0210-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: HEPA DNA VIRUS RECEPTOR NUMBER OF SEQUENCES: 117
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                            TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                    LENGTH: 55 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatil
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RESULT 2
US-08-480-173A-44
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GENERAL INFORMATION:
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Best Local Similarity 100.0%;
Matches 12; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                         Sequence 44, Application US/08484408A Patent No. 6117653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPAX: 612-334-8994 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Thoma, Hans A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALCHIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,173A
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Thoma, Hans A
TITLE OF INVENTION: HEPATITIS B SURFACE ANTIGEN VACCINE
NUMBER OF SEQUENCES: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 174 amino acids
TYPE: amino acid
STRANDENNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: PODOVICH, Thomas E
REGISTRATION NUMBER: 30,099
REFERENCE DOCKET NUMBER: MED1003USD4
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-334-8991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/484,408A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: ur
                                                                                                                                                                                                                                                           TITLE OF INVENTION: HEPATITIS B SURFACE ANTIGEN VACCINE NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     160 PLSSIFSRIGDP 171
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CITY: Minneapolis
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                                                                                                                                                                                    STREET: 80 S. 8th CITY: Minneapolis STATE: MN
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                                                                                                                                               COUNTRY: UZIP: 55402
                                                                                                                                                                                                                                               ADDRESSEE:
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80 S. 8th Street, Suite 1902
                                                                                                                                                                                                                          E: Popovich & Wiles, P.A.
80 S. 8th Street, Suite 1902
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                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 214, Application US/08105483 Patent No. 5494807
                                                                                                                                                                                   COMPUTER: INFOCUTION COMPACIBLE
COMPUTER: INFOCUTION (MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PACENTIN Release #1.0, Version #1.25
SOPTWARE: PACENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/105,483
FILING DATE: 12-AUG-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/847,951
FILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2400
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEPHONE: (212) 840-333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 612-334-8994
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION: APPLICANT: Paolet
Query Match
Best Local Similarity
                                                                                                                                                         TELEFAX: (212) 840-0712 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Popovich, Thomas E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: un
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 160 PLSSIFSRIGDP 171
                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 281 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: New York
STATE: NY
COUNTRY: USA
ZIP: 10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 30,099
REFERENCE/DOCKET NUMBER: MEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                                                                     STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                174 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      530 Fifth Avenue
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                                                                ESS: single
linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unknown
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 100.0%;
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   Score 61; DB 1; Pred. No. 0.0006;
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                    Length 281;
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RESULT 6
US-08-458-101-214
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                                                                                                                                                                                                                               Sequence 21
Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%; S
Best Local Similarity 100.0%; p
                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 5762938
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                              APPLICANT:
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MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 21-AUG-19
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ADDRESSEE: Curtis, Morris & Safford
ADDRESSEE: C/o William S. Frommer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                        41 PLSSIFSRIGDP 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 281 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                1 PLSSIFSRIGDP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 10036
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o. 5766599
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                                                               No. 57665
Riviere,
                                                                  Tartaglia, James
No. 5766599ton, Elizabeth K.
Riviere, Michel
                                                                                                                                    Perkus, Marion E.
Taylor, Jill
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                                                                                                                                                                                   Paoletti, Enzo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patentin Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ) 840-0712
NO: 214:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 61; DB 1;
Pred. No. 0.0006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 281;
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                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 216, Application
Patent No. 5494807
GENERAL INFORMATION:
APPLICANT: Pacoletti,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/458,101
FILING DATS: 01-UW-1995
CLASSIFICATION: 424
ATTORNEY/AGENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-BOS/MS-DOS
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TELEBEAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: GENETIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 25,506
TELECOMMUNICATION INFORMATION:
TELEPHONE: /2707
                                                                                                                                                                                                         STREET: 530 Fift
CITY: New York
STATE: NY
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NAME: Frommer, William S
REGISTRATION NUMBER: 25,
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APPLICANT: Johnson, Gerard P.
APPLICANT: Pincus, Steven E.
APPLICANT: Cox, William I.
APPLICANT: Gettig, Russell Robert
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN
NUMBER OF SEQUENCES: 467
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APPLICANT:
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ADDRESSEE: c/o William S. Frommer
STREET: 530 Fifth Avenue
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TOPOLOGY: 1i
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ADDRESSEE: c/o William S. Frommer
STREET: 530 Fifth Avenue
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), 5494807
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                                                                                                                                                                                        USA
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                                                                                                                                                                                                                                                                                                                                                  462
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Pred. No. 0.0006;
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CURRENT APPLICATION DATA:

CLASSIFICATION: APPLICATION NUMBER:

12-AUG-1993

US/08/105,483

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NAME: Prommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 4543:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-333
TELEPHONE: (212) 840-333
TELEPHAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 219:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
TYPE: amino acids
TYPE: amino acids
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 5494807
GENERAL INFORMATION:
APPLICANT: Paolet
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REFERENCE/DOCKET NUMBER: 45'
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEPAX: (212) 840-0712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/847,951
APPLICATION NUMBER: US 07/847,951
FILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: POPPY disk
COMPUTER: IBM PC compatible
COMPUTER: PC COMPATION:
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/105,483
FILING DATE: 12-AUG-1993
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 462
CORRESPONDENCE ADDRESS:
CORTESPONDENCE CUrtis, Morris & Safford ADDRESSEE: C/o William S. Frommer STREET: 530 Fifth Avenue
                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/847,951
PILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
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CITY: New York
STATE: NY
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                                                                              RESULT 10
US-08-709-209-219
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Best Local Similarity
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                                                                                                                                                                                                        Query Match
Best Local S
Matches 12
                                             Sequence 219, Application US/08709209 Patent No. 5762938
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/709,209
APPLICATION = 21-AUG-1996
CLASSIFICATION = 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/105,483
APPLICATION NUMBER: US 07/847,951
APPLICATION NUMBER: US 07/847,951
FILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Paolet
GENERAL INFORMATION:
APPLICANT: Paoletti, Enzo
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: (
TITLE OF INVENTION: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: c/o William S.
STREET: 530 Fifth Avenue
CITY: New York
                                                                                                                                             149 PLSSIFSRIGDP 160
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                                                                                                                                                                                               100.0%;
Local Similarity 100.0%;
nes 12; Conservative
                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2400
                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX:
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                                                                                                                                                                                                                                                                                                                                       ENGTH:
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                                                                                                                                                                                                                                                                                        linear
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Pred. No. 0.00086;
                                                                                                                                                                                                                             Score 61; DB 1;
Pred. No. 0.00086;
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US-08-458-101-216
; Sequence 216, Application US/08458101
; Patent No. 5786599
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/847,
FILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 4543
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/
FILING DATE: 21-AUG-1996
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 12-AUG-1932
CORRESPONDENCE ADDRESS:
ADDRESSEE: C/O William S. Frommer
STREET: 530 Fifth Avenue
                                                                                                                         APPLICANT:
                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                               APPLICANT:
                                                                 APPLICANT: Audonnet, Jean-Christophe Francis
APPLICANT: Gettig, Russell Robert
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN
MUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: JEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 389 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 PLSSIFSRIGDP 12
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CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CENGTH:
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ADDRESSEE: c/o William S. Frommer
STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OF SEQUENCES:
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Limbach, Keith J.
Johnson, Gerard P.
Pincus, Steven E.
Cox, William I.
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No. 5766599con, Elizabeth K.
Riviere, Michiga
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Taylor, Jill
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linear
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US-08-458-101-219
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TELEPAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 216:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
TYPE: amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-458-101-216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/458,101
FILING DATE: 01-JUN-1995
CLASSIFICATION: 424
ATTORNEY/AGRAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/458,10:
                                                                                                                                                                                                                                                                  APPLICANT: Limbach, Keith J.
APPLICANT: Johnson, Gerard P.
APPLICANT: Pincus, Steven B.
APPLICANT: Cox, William I.
APPLICANT: Audonnet, Jean-Christophe Francis
APPLICANT: Gettig, Russell Robert
TITLE OF INVENTION: GENETICALLY ENGINEERED VA
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: C/O William S. Frommer
STREET: 530 Fifth Avenue
                                                                                                                                                   STREET: 530 Fif
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2740
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                    COUNTRY:
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TOPOLOGY: lir
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STATE: NY
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                                                                                                                                     USA
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No. 5766599ton, Elizabeth
Riviere, Michel
Riviere, Charles
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                                                                                                                                                                                                                                                                    GENETICALLY ENGINEERED VACCINE STRAIN
                                                                                                                                                                                                                                                      467
US/08/458,101
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Pred. No. 0.00086;
; Mismatches 0;
                                 Version #1.25
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                                                                                                                                                                                                                                                                                      ZIP: 10036-2711

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/486,099

FILING DATE: 07-JUN-1995

CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                      TELEFAX: (212) 869-9741/886/
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: lin-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF
TITLE OF INVENTION: MEMBRANE FUSION-ASSOCIATED EVENTS, INCLUDING HEPATITIS
TITLE OF INVENTION: B VIRUS TRANSMISSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: COCUZZÍ, LANIE A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-031
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   149 PLSSIFSRIGDP 160
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CITY: New York
STATE: New York
COUNTRY: USA
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REFERENCE/DOCKET NUMBER: 454310-2740
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CLASSIFICATION:
WOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
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                                      STRANDEDNESS
                                                                          ENGTH:
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o. 6013263
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                                                      amino acids
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1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bolognesi, Dani I
Matthews, Thomas
Wild, Carl T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Barney, Shawn O
                                                                                                                                                     (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lambert, Dennis M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-1995
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Thomas J.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/360,107A
FILING DATE: 20-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COPUZZI, LAURA A.
NAME: COPUZZI, LAURA A.
                                                                                                                Query Match
                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                    TELEFAX: (212) 869-9741.
TELEX: 66141 PENNIE
TRIEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION
TITLE OF INVENTION: OF MEMBRANE FUSION-ASSOCIATED EVENTS, INTITLE OF INVENTION: TRANSMISSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                    NAME: Coruzzi Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7877
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 149 PLSSIFSRIGDP 160
                                                                                                                                                                          TOPOLOGY: unknown MOLECULE TYPE: protein
  149 PLSSIFSRIGDP 160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PLSSIFSRIGDP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                       / Match 100.0%; Score 61; DB 3; Local Similarity 100.0%; Pred. No. 0.00086; Local Similarity 0, Mismatches 0;
                                                                                                                                                                                                               STRANDEDNESS:
                                    1 PLSSIFSRIGDP 12
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                                                                                                                                                                                                                                    amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bolognesi, Dani P.
Matthews, Thomas J.
Wild, Carl T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Barney, Shawn O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lambert, Dennis M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 61; DB 3; 1
100.0%; Pred. No. 0.00086;
ative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                    7872-013
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                                                                                                                       Length 389;
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                                                                                    Indels
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                                                                                      Gaps
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RESULT 15 US-08-484-223B-106 ; Sequence 106, Application US/08484223B ; Patent No. 6020459

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GENERAL INFORMATION:

APPLICANT: Botthews, Thomas J.

APPLICANT: Matthews, Thomas J.

APPLICANT: Matthews, Thomas J.

APPLICANT: Hathews, Thomas J.

APPLICANT: Hathews, Thomas J.

APPLICANT: Hathews, Thomas J.

APPLICANT: Hathews, Thomas J.

APPLICANT: Hathert, Dennis M.

COMMUTER: Hathert, Dennis M.

ACTIT: New York

COUNTRY: USA,

ZIF: 1003-3711

COUNTRY: USA,

ZIF: 1003-3711

COUNTRY: USA,

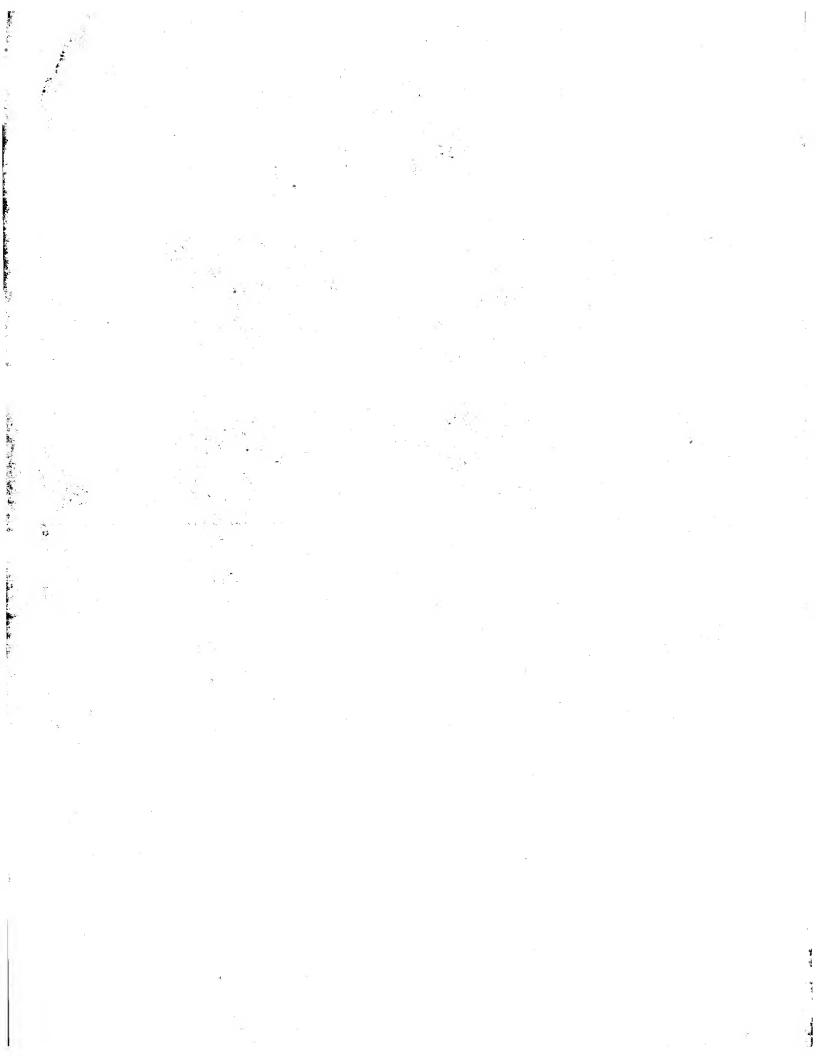
ZIF: 1004-3711

COUNTRY: HATHERT M.

COUNTRY: HATHERT M.

APPLICANTION M.

APPLICANTION
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-890-752A-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            122226 seqs, 20178551 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 PLSSIFSRIGDP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     February 3, 2003, 09:25:53; Search time 0.439024 Seconds (without alignments) 551.547 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Published_Applications_AA: *
/cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US00_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US00_NEW_PUB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US00_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES	

Result	Score	Query	Query Match Length	DB	ID	Description
ب	61	100.0	281	ا و	US-09-247-890-12	Semience 10 Ann
N	59	96.7	174	10	US-09-818-066-55	т п
L	59	96.7	389	9	US-09-821-877-2	۲.
4	54		170	10	US-09-818-066-49	٠,
₅	54	88.5	174	10	US-09-818-066-46	4
o	54		174	5	US-09-818-066-48	
7	54	88.5	174	10	US-09-818-066-56	Sequence 56 App
œ	54	88.5	281	ø	US-09-247-890-10	10. A
9	54		400	ø	US-10-209-264-3	Sequence 3. Appl
10	50		174	10	US-09-818-066-50	٠,
11	52		174	10	US-09-818-066-58	
12	49		174	10	US-09-818-066-53	J .
13	46		174	10	US-09-818-066-45	
14	46	75.4	174	10	US-09-818-066-47	47
15	44	72.1	174	10	US-09-818-066-37	37
16	44	72.1	174	10	US-09-818-066-54	Sequence 54. App.
17	43	70.5	174	10	US-09-818-066-57	7
18	40	65.6	64	10	US-09-917-340-85	Sequence 85. Appl
19	40	עעעע	174	5	75-09-818-066-43))

45	44	43	42	41	40	39	38	37	36	35	34	33	32	4	0	29	28	27	26	25	24	23	22	21	20
3.4	34 4	34	ω 4.	ų A	LJ AL	ىن 4.	35	35	35	35	<u>3</u> 5	35	35	35	35	S	36	36	36	36	37	38	38	38	40
55.7	55.7	55.7	55.7	55.7	55.7	55.7	57.4	57.4	•	57.4	57.4		57.4	57.4	57.4	57.4				59.0	60.7	62.3	62.3	62.3	65.6
478	312	284	174	174	119	64	1036	1036	743	743	705	703	690	174	173	96	174	174	174	55	174	174	174	174	174
9	10	ø	10	10	10	10	10	10	10	10	9	5	5	10	ø	5	10	10	10	10	10	10	10	10	10
US-10-156-634A-2	US-09-801-368-260	US-09-895-913A-102	US-09-818-066-44	US-09-818-066-43	US-09-925-299-799	US-09-917-340-84	US-09-771-161A-256	US-09-771-161A-255	US-09-771-161A-254	US-09-771-161A-164	US-09-738-626-4055	US-09-815-242-5610	US-09-815-242-12460	US-09-818-066-39	US-10-051-643-207	US-09-864-761-39796	US-09-818-066-52	US-09-818-066-36	US-09-818-066-35	US-09-879-257A-45	US-09-818-066-59	US-09-818-066-41		US-09-818-066-38	US-09-818-066-62
Sequence 2, Appli	Sequence 260. App	102	44	43,	799	84,		255	254	164	4055		Sequence 12460. A	w	Sequence 207, App	3979	52	U O		45	59	41,	40	38,	Sequence 62, Appl

ALIGNMENTS

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RESULT 2
US-09-818-066-55
US-09-818-066-55; Sequence 55, Application US/09818066; Patent No. US20020032307A1
GENERAL INFORMATION:
APPLICANT: Shuping Tong et al.
                                                                                                                                                                                                                                                                                                                                                                                      US-09-247-890-12

Sequence 12, Application US/09247890

Publication No. US20020198162A1

GENERAL INFORMATION:

APPLICANT: Punnonen, Juha
APPLICANT: Bass, Steven H.

APPLICANT: Whalen, Robert Gerald
APPLICANT: Whalen, Robert H.

APPLICANT: Maxygen, Inc.

FITLE OF INVENTION: Antigen Library Immunization
FILE REFERENCE: 018097-028710US

CURRENT FILING DATE: 1999-02-10

FEARLIER APPLICATION NUMBER: US/09/247,890

CURRENT FILING DATE: 1998-02-11

FEARLIER APPLICATION NUMBER: US 60/074,294

EARLIER APPLICATION NUMBER: US 60/074,294

FEARLIER FILING DATE: 1998-02-11

FEARLIER PRICHED DATE: 1998-10-23

NUMBER OF SEQ ID NOS: 25

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 12

LENGTH: 281

TYPE: PAT

ORGANISM: Hepatitis B virus

US-09-247-890-12
                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Watches 12; Conserve
                                                                                                                                                                                                                       밁
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                                                                                                                                                                                                                    41 PLSSIFSRIGDP 52
                                                                                                                                                                                                                                                     1 PLSSIFSRIGDP 12
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                                                                                                                                                                                                                                                                                                                                               Length 281;
                                                                                                                                                                                                                                                                                                        Indels
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0

Gaps

TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS:

ADDRESSEE: Fish & Richardson P.C.

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US-09-821-877-2
                                                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 389
TYPE: PRT
ORGANISM: Hepatitis B Virus
US-09-821-877-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-818-066-55
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09821877 Patent No. US20020177124A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 91.7
Matches 11; Conservative
                                                Query Match
Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                              APPLICANT: Abbott Laboratories
APPLICANT: Coleman, Paul F.
APPLICANT: Mushahwar, Isa K.
TITLE OF INVENTION: Hepatitis B Virus Surface Antigen Mutant
TITLE OF INVENTION: And Methods Of Detection Thereof
FILE REFERENCE: 6794.US.O1
CURRENT APPLICATION NUMBER: .US/09/821,877
CURRENT FILING DATE: 2000-03-03
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 55: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Ploppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/818,066
PILING DATE: 27-Mar-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:

CLASSIFICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             160 PISSIFSRIGDP 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 PLSSIFSRIGDP 12
1 PLSSIFSRIGDP 12
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MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 55:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/683,262 FILING DATE: 18-JUL-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 174 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX:
                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02110-2804
                                                                  96.7%;
91.7%;
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91.7%; Pred. No. 0.00073;
ative 1; Mismatches 0
                                     Score 59; DB 9; Pred. No. 0.0017; Indels
                                                  1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               00786/287002
                                                                                         DB 9; Length 389;
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TOPOLOGY: linear;
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 49:
US-09-818-066-49
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                                                                                                                                                                                                                                                                                                                RESULT 5
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Patent No. US20020032307A1
GENERAL INFORMATION:
                                                                                                                 Sequence 46, Application US/09818066
Patent No. US20020032307A1
GENERAL INFORMATION:
APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 200154
INFORMATION FOR SEQ ID NO: 49:
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APPLICATION NUMBER: US 08/683,262
FILING DATE: 18-JUL-1996
ATTORNEY/AGENT INFORWATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE/DOCKET NUMBER: 00786/287002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                               156 PISSIFSRIGDP 167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/818,066
FILING DATE: 27-Mar-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |:|||||||||
|PISSIFSRIGDP 160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                      PLSSIFSRIGDP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Boston
                                     STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 170 amino acids TYPE: amino acid
                     ZIP: 02110-2804
                                                                                CITY: Boston
                                                                                                    STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 54; DB 10; Length 170;
Pred. No. 0.0058;
1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 48, Application upatient No. US20020032307A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
    INFORMATION FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: 200154
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
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160 PISSIFSRTGDP 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 88.5%;
Local Similarity 83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PLSSIFSRIGDP 12
                                                    NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE/DOCKET NUMBER: 00786/287002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/683,262
FILING DATE: 18-JUL-1996
                                                                                                                                           ATTORNBY/AGENT INFORMATION:
NAME: Fraser, Janis K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE/DOCKET NUMBER: 007
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/683,262 FILING DATE: 18-JUL-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE DESCRIPTION: SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/818,066
FILING DATE: 27-Mar-2001
CLASSIFICATION: <Unknown>
                                 TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 174 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/818,066 FILING DATE: 27-Mar-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WDDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/09818066
                200154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         225 Franklin Street
ID NO: 48:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
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Pred. No. 0.006;
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US-09-818-066-56
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                                                                                         Best
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Best Local Similarity
Matches 10; Conserv
                                                                        Matches
                                                                                                           Query Match
                                                                                                                                                                                                                                                    TELEX: 200154
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/818,066
FILING DATE: 27-Mar-2001
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION UMBER: US 08/683,262
FILING DATE: 18-JUL-1996
ATTORNEY/AGENT IMPORMATION:
NAME: Praser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE/DOCKET NUMBER: 00786/287002
TELECOMUNICATION INFORMATION:
TRUIDENCH INFORMATION:
TRUIDENCH INFORMATION:
TRUIDENCH INFORMATION:
TRUIDENCH INFORMATION:
TRUIDENCH INFORMATION:
TRUIDENCH INFORMATION:
160
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                                                                                       Local Similarity
                                1 PLSSIFSRIGD 11
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                                                                                                                                                        MOLECULE TYPE: protein SEQ ID NO: 56:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
NUMBER OF SEQUENCES: 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 48:
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PLSSIFSRIGD 170
                                                                    11;
                                                                                                                                                                                                  TOPOLOGY: linear
                                                                                                                                                                                                                                 LENGTH: 174 amino acids
                                                                                                                                                                                                                                                                                                        TELEFAX:
                                                                                                                                                                                                                                                                                                                               TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Massachusetts
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                                                                      Conservative
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                                                                                                                                                                                                                    amino acid
                                                                                                                                                                                                                                                                                                          (617) 542-8906
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                                                                                    88.5%;
100.0%;
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                                                                  Score 54; DB 10;
Pred. No. 0.006;
0; Mismatches 0;
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Pred. No. 0.006;
                                                                                                     Length 174;
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                                                                  Indels
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RESULT 8 US-09-247-890-10 ; Sequence 10, Application US/09247890

Publication No.

US20020198162A1

INFORMATION:

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APPLICANT: Maxygen, Inc.
TITLE OF INVENTION: Antigen Library Immunization
FILE REFERENCE: 018097-028710US
CURRENT APPLICATION NUMBER: US/09/247,890
CURRENT FILLING DATE: 1999-02-10
EARLIER APPLICATION NUMBER: US 60/074,294
EARLIER FILLING DATE: 1998-02-11
EARLIER FILLING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 281
TYPE: PAT
ORGANISM: Hepatit18 B virus
US-09-247-890-10
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US-10-209-264-3
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Publication No. US20030003111A1
GENERAL INFORMATION:
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Best Local &
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APPLICANT: Bass, Steven H.
APPLICANT: Whalen, Robert Gerald
APPLICANT: Howard, Russell
APPLICANT: Stemmer, Willem P.C.
                                       TELEPHONE: (212) 70 INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41 PISSIFSRTGDP 52
                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/SG98/00046

FILING DATE: 19-JAN-1998

ATTORNEY/AGENT INFORMATION:

NAME: Mass, Clifford J.

REGISTRATION NUMBER: 30,086

REFERENCE/DOCKET NUMBER: U-013109-7

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 88.5%;
Local Similarity 83.3%;
es 10; Conservative
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                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBP C COMPATIBLE

COMPUTER: PLOS C COMPATIBLE

COMPUTER: PROCEDOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zhao, Yi
Chen, Wei Ning
TITLE OF INVENTION: A MUTANT HUMAN HEPATITIS B VIRAL STRAIN AND
USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Oon, Chong Jin
Lim, Gek Keow
                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/10/209,264
PILING DATE: 31-Jul-2002
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Ladas & Parry
STREET: 26 West 61 Street
LENGTH: 400 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                  10023
                                                               708-1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Mismatches
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Pred. No. 0.01;
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RESULT 11
US-09-818-066-58
US-09-818-066-58
; Sequence 58, Application US/09818066
; Patent No. US20020032307A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: protein ; SEQUENCE DESCRIPTION: SEQ ID NO: 50: US-09-818-066-50
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Best Local Similarity 83.3%;
Matches 10; Conservative
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GENERAL INFORMATION:
                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                           TELEX: 200154
INFORMATION FOR SEQ ID NO: 50:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION UNMEER: US/99/818,066
FILING DATE: 27-Mar-2001
CLASSIFICATION: JUNKnown>
PRIOR APPLICATION NUMBER: US 08/683,262
FILING DATE: 18-JUL-1996
ATTORNEY/AGENT INFORMATION:
NUMBER: FYSTER: JUNIER
                                                                                                                                  161 IŚŚIFŚRIGDP 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 PLSSIFSRIGDP 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A. ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                           LENGTH: 174 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/09818066
                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 225 Franklin Street
                                                                                                                                                                                                                       85.2%;
90.9%;
                                                                                                                                                                                                   Score 52; DB 10; pred. No. 0.014; 1; Mismatches
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Pred. No. 0.015;
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RESULT 12
US-09-818-066-53
/ Sequence 53, Application US/09818066
/ Patent No. US20020032307A1
/ GENERAL INFORMATION:
/ APPLICANT: Shuping Tong et al.
/ TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
/ NUMBER OF SEQUENCES: 75
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; SEQUENCE DESCRIPTION: SEQ ID NO: 58:
US-09-818-066-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 10; Conserv
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INFORMATION FOR SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/09/818,066
FILING DATE: 27-Mar-2001
CLASSIFICATION: CUnknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        161 ISSIFSRIGDP 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 LSSIFSRIGDP 12
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ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Praser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE DOCKET NUMBER: 007
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/683,262
FILING DATE: 18-JUL-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOPTMARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION UNMER: US/09/818,066
FILING DATE: 27-Mar-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
                                                                                                                                                                                                                                   STATE: Massachusetts
                                                                                                                                                                                                                                                       CITY: Boston
                                                                                                                                                                                             ZIP: 02110-2804
                                                                                                                                                                                                                                                                        STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 174 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: (617) 542-50
(617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  85.2%;
90.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 52; DB 10; Length 174; Pred. No. 0.014;
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US-09-818-066-45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 81.8
Matches 9; Conservative
                                                                                                              TELEX: 200154
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 200154
INFORMATION FOR SEQ ID NO: 53:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PATENTIN REJEASE #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/818,066 FILING DATE: 27-Mar-2001 CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
        TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 45:
                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 00786/287002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE/DOCKET NUMBER: 00'
TELECOMMUNICATION INFORMATION:
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FILING DATE: 18-JUL-1996
ATTORNEY/AGENT INFORMATION:
                                                                    LENGTH: 174 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/683,262 FILING DATE: 18-JUL-1996
                                                                                                                                                                        TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Massachusetts
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (617) 542-8906
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                80.3%;
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Pred. No. 0.049;
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Query Match

75.4%; Score 46;

DB 10;

Length 174;

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US-09-818-066-47
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US-09-818-066-37
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Patent No. US20020032307A1
GENERAL INFORMATION:
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               Sequence 37, Application US/09818066
Patent No. US20020032307A1
GENERAL INFORMATION:
APPLICANT: Shuping Tong et al.
APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 75.0
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617) 542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 47:
SEQUENCE CHARACTERISTICS:
LENGTH: 174 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            160 PISSISSRTGDP 171
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FILING DATE: 27-Max-2001
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATE: US 08/683,262
APPLICATION NUMBER: US 08/683,262
FILING DATE: 18-JUL-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                         160 PISSISSRTGDP 171
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PARENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR
NUMBER OF SEQUENCES: 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE/DOCKET NUMBER: 00786/287002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Pish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 47:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Boston
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                            75.4%; Score 46; DB 10; Length 174; 75.0%; Pred. No. 0.18;
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1; Mismatches

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Query Match
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Search completed: February
Job time : 0.439024 secs
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INFORMATION FOR SEQ ID NO: 37:
SEQUENCE CHARACTERISTICS:
                                                                              161 ISSISSRIGDP 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/818,066
FILING DATE: 27-MAX-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/683,262
FILING DATE: 18-JUL-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                   2 LSSIFSRIGDP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 02110-2804
COMPUTER READABLE FORM:
COMPUTER: Floppy disk
COMPUTER: IBM FC COMPATIBLE
COMPUTER: FOR FC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                      MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 37:
                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: (
TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                             LENGTH: 174 amino acida
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (617) 542-8906
                                                                                                                                                         Conservative
                                                                                                                                                                             72.1%;
81.8%;
                      3, 2003, 09:35:29
                                                                                                                                                           1; Mismatches
                                                                                                                                                                             Score 44; DB 10; Length 174;
Pred. No. 0.41;
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Minimum DB
Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Searched:
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Perfect score:
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                           protein +
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length: 2000000000
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1: /cgn2_6/ptodata/1/paa/PCT NEW COMB.pep:*

2: /cgn2_6/ptodata/1/paa/US06_NEW COMB.pep:*

3: /cgn2_6/ptodata/1/paa/US07_NEW COMB.pep:*

4: /cgn2_6/ptodata/1/paa/US07_NEW COMB.pep:*

5: /cgn2_6/ptodata/1/paa/US09_NEW COMB.pep:*

6: /cgn2_6/ptodata/1/paa/US09_NEW COMB.pep:*

7: /cgn2_6/ptodata/1/paa/US00_NEW COMB.pep:*
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                         BLOSUM62
Gapop 10.0 , Gapext 0.5
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Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Copyright
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2003, 09:24:27; Search time 0.982578 Seconds (Without alignments) 988.252 Million cell updates/sec
             US-10-211-088-302
US-10-326-908-19
US-10-326-908-19
US-10-326-908-21
PCT-US02-2263-46
US-10-287-994-46
PCT-US02-40225-314
US-10-320-797-3144
US-10-320-797-3144
US-10-092-411A-4053
PCT-US02-32727-2247
PCT-US02-32727-1919
US-10-057-498-2247
PCT-US02-32727-23079
US-10-057-498-23079
US-10-057-498-23079
US-10-057-498-23079
US-10-218-140-3758
US-09-724-676-70237
US-09-724-676-70238
US-09-724-676-70238
PCT-US02-34321-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUMMARIES
Sequence 302, App Sequence 21, Appl Sequence 21, Appl Sequence 46, Appl Sequence 23, Appl Sequence 3144, Ap Sequence 3144, Ap Sequence 2247, Ap Sequence 2247, Ap Sequence 23079, A Sequence 23079, A Sequence 70237, A Sequence 70237, A Sequence 70237, A Sequence 70238, A Sequence 70238, A Sequence 70231, A Sequence 70230, A Sequence 70231, A
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                                                  Sequence 19, Application US/10326908

GENERAL INFORMATION:
APPLICANT: Christopher JONES
APPLICANT: Ghistopher JONES
APPLICANT: Gill DOUCE
APPLICANT: Gill DOUCE
APPLICANT: Mark PAGE
TITLE OF INVENTION: Designing Immunogens
FILE REFERENCE: 117-433 / N78027D
CURRENT APPLICATION NUMBER: US/10/326,908
CURRENT FILING DATE: 2007-1210-036
                                                                                                                                                                                                                                                                                      US-10-326-908-19
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US-10-211-088-302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local
Matches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                    PRIOR APPLICATION NUMBER: US 10/110,036 PRIOR FILING DATE: 2002-04-08
    APPLICATION NUMBER:
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US-09-724-676-70225	US-09-724-676-70224	US-09-724-676A-70241	US-US-124-6/6A-/U240	110-00-011 CTC TOTAL	US-09-724-676-70241	US-09-724-676-70240	US-09-724-676A-70239	US-09-724-676A-70234	US-09-724-676-70239	00-09-724-6/6-70234	TO DO TO	US-09-724-676A-70223	US-09-724-676-70223	US-10-057-498-1231	PCT-US02-32727+1231	US-US-124-6/6A-70233		US-09-724-676A-70232	US-09-724-676-70233	00 00 7EH: 070 70EGA	IIS-09-704-676-70000	US-09-724-676A-70231
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70225,	70224	70241, /	70240, 1	/0241, /	1000	70240	70239	70234	70239,	70234,	10223, 1	30000	~	1231. An	1231, A	70233,	, ,,,,,,	3000	70233	10232,	1000	70031
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ALIGNMENTS

Sequence 302, Application US/10211088 GENERAL INFORMATION: APPLICANT: Bright, Gary R. APPLICANT: Premkumar, D. David APPLICANT: Chen, Yih-Tai TITLE OF INVENTION: Novel Fusion Proteins And Assays For Molecular Binding FILE REFERENCE: 01-1022-US SOFTWARE: PatentIn version 3.1 SEQ ID NO 302 LENGTH: 12 CURRENT APPLICATION NUMBER: US/10/211,088 CURRENT FILING DATE: 2002-10-15 PRIOR APPLICATION NUMBER: 60/309,395 PRIOR FILING DATE: 2001-08-01 PRIOR APPLICATION NUMBER: 60/341,589 PRIOR FILING DATE: 2001-12-13 NUMBER OF SEQ ID NOS: 366 TYPE: PRT ORGANISM: Artificial sequence FEATURE: OTHER INFORMATION: Protein-derived transport 1 PLSSIFSRIGDP 12 100.0%; |llarity 100.0%; |Conservative 0; 0 Score 61; DB 6; 1 Pred. No. 4.6e-05; Mismatches 0; peptide Length 12; Indels 0 Gaps 0

PCT/GB00/03857

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PRIOR FILING DATE: 2000-10-06

PRIOR APPLICATION NUMBER: GB 0007789.1

PRIOR APPLICATION NUMBER: GB 9007789.1

PRIOR APPLICATION NUMBER: GB 9923902.2

PRIOR FILING DATE: 1999-10-08

NUMBER OF SEQ ID NOS: 30

SOFTWARE: MS WORD

SEQ ID NO 19

LENGTH: 64

TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: synthetic peptide immunogen
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CURRENT APPLICATION NUMBER: US/10/326,908
CURRENT FILING DATE: 2002-12-23
PRIOR APPLICATION NUMBER: US 10/110,036
PRIOR PILING DATE: 2002-04-08
PRIOR PPLICATION NUMBER: US 10/110,0387
PRIOR PILING DATE: 2000-10-06
PRIOR PILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: GB 0007789.1
PRIOR APPLICATION NUMBER: GB 9923902.2
PRIOR PILING DATE: 2000-03-30
PRIOR PILING DATE: 1999-10-08
PRIOR PILING DATE: 1999-10-08
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                                                                                                                                             RESULT 4
PCT-US02-32263-46
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Best Local Similarity 91.7
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 21
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                                                                                    Sequence 46, Application PC/TUS0232263
GENERAL INFORMATION:
APPLICANT: Neose Technologies, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Christopher JONES
APPLICANT: Andrew BACON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Designing Immunogens FILE REFERENCE: 117-433 / N78027D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 30
SOFTWARE: MS Word
                                                   APPLICANT:
                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           48 PISSIPSRIGDP 59
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Mark PAGE
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Bayer, Robert
Bowe, Caryn
Hakes, David
                                                 Zopf, David
                                                                        DeFrees, Shawn
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91.7%;
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Pred. No. 0.00061
1; Mismatches
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Pred. No. 0.00064;
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PRIOR APPLICATION NUMBER: US 60/328,523
PRIOR FILING DATE: 2001-10-10
PRIOR APPLICATION NUMBER: US 60/344,692
PRIOR FILING DATE: 2001-10-19
PRIOR APPLICATION NUMBER: US 60/344,692
PRIOR APPLICATION NUMBER: US 60/334,233
PRIOR APPLICATION NUMBER: US 60/334,301
PRIOR FILING DATE: 2001-11-08
PRIOR FILING DATE: 2001-11-08
PRIOR PILING DATE: 2002-06-07
PRIOR APPLICATION NUMBER: US 60/387,292
PRIOR APPLICATION NUMBER: US 60/387,777
PRIOR APPLICATION NUMBER: US 60/391,777
PRIOR APPLICATION NUMBER: US 60/391,777
PRIOR FILING DATE: 2002-06-25
PRIOR FILING DATE: 2002-06-25
PRIOR FILING DATE: 2002-08-16
PRIOR APPLICATION NUMBER: US 60/404,249
PRIOR FILING DATE: 2002-08-16
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PCT-US02-32263-46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 5
                                                                       PRIOR APPLICATION NUMBER: US 60/391,777
PRIOR FILING DATE: 2002-06-25
PRIOR APPLICATION NUMBER: US 60/396,594
PRIOR APPLICATION NUMBER: US 60/404,249
PRIOR APPLICATION NUMBER: US 60/404,249
PRIOR FILING DATE: 2002-08-16
PRIOR APPLICATION NUMBER: US 60/407,527
PRIOR APPLICATION NUMBER: US 60/407,527
PRIOR FILING DATE: 2002-08-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.1 SEQ ID NO 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 46, Application US/10287994 GENERAL INFORMATION:
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Best Local Similarity
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NUMBER OF SEQ ID NOS: 62
SOFTWARE: PatentIn version 3.1
SEQ ID NO 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2002-08-28
NUMBER OF SEQ ID NOS: 62
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TYPE: PRT
                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: REMODELING AND GLYCOCONJUGATION OF PEPTIDES
FILE REFERENCE: 040853-01-5052-00
CURRENT APPLICATION NUMBER: US/10/287,994
CURRENT FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: US 60/328,523
PRIOR FILING DATE: 2001-10-10
PRIOR APPLICATION NUMBER: US 60/344,692
PRIOR FILING DATE: 2001-10-19
PRIOR FILING DATE: 2002-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Neose Technologies, APPLICANT: DeFrees, Shawn
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Chen, Xi
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Bowe, Caryn
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Pred. No. 0.036;
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APPLICANT: Paul, Ralph
APPLICANT: Cudmore, Sally
APPLICANT: Cudmore, Sally
APPLICANT: Cudmore, Sally
APPLICANT: Cudmore, Daniel J.
TITLE OF INVENTION: LIPID-COMPRISING DRUG DELIVERY COMPLEXES
TITLE OF INVENTION: AND METHODS FOR THEIR PRODUCTION
FILE REFERENCE: 226272005340
CURRENT APPLICATION NUMBER: PCT/US02/13609
CURRENT PILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/287,786
PRIOR FILING DATE: 2001-04-30
NUMBER OF SEQ ID NOS: 45
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 23
LENGTH: 10
TYPE: PRT
COGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                        RESULT 7
PCT-US02-40225-3144
; Sequence 3144, Application PC/TUS0240225
; GENERAL INFORMATION:
                                                             PCT-US02-40225-3144
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; OTHER INFORMATION: Synthetic Construct
PCT-US02-13609-23
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                                                                                                                               APPLICANT: Elitra Pharmaceuticals, Inc.
APPLICANT: Eroshkin, Alexey M.
APPLICANT: Eroshkin, Alexey M.
APPLICANT: Zamudio, Carlos
TITLE OF INVENTION: IDENTIFICATION OF ESSENTIAL GENES OF CRYPTOCCCCUS NEOFORMANS AND
TITLE OF INVENTION: METHODS OF USE
FILE REFERENCE: 10182-021-228
CURRENT APPLICATION NUMBER: PCT/US02/40225
CURRENT FILING DATE: 2002-12-17
PRIOR APPLICATION NUMBER: 60/341,261
PRIOR PILING DATE: 2001-12-17
NUMBER OF SEQ ID NOS: 3361
SOPTWARE: PatentIn version 3.1
SEQ ID NO 3144
  Query Match
Best Local Similarity
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Matches 10; Conservative
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                                                                            ORGANISM: Cryptococcus neoformans
                                                                                                    TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -US02-13609-23
                                                                                                                       LENGTH: 399
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  63.9%;
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Score 39;
Pred. No.
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Pred. No. 0.036;
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DB 1;
25;
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CURRENT APPLICATION NUMBER: US/10/320,797
CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: 60/341,261
PRIOR FILING DATE: 2001-12-17
NUMBER OF SEQ ID NOS: 3361
SOPTWARE: Patentin version 3.1
SEQ ID NO 3144
LENGTH: 399
TYPE: PRT
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                                     Ş
                                                                                                                                               ; ORGANISM: Staphylococcus epidermidis
US-10-092-411A-4053
                                                                                                                                                                                                                                                  PILE REFERENCE: 032796-101
CURRENT APPLICATION NUMBER: US/10/092,411A
CURRENT FILING DATE: 2002-03-07
PRIOR APPLICATION NUMBER: US 09/134,001
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Cryptococcus neoformans US-10-320-797-3144
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US-10-320-797-3144
                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 5676
SEQ ID NO 4053
LENGTH: 533
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4053, Application US/10092411A
GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                         Matches
                                                                                          Query Match
Best Local
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GENERAL INFORMATION:
APPLICANT: Eroshkin, Alexey M.
APPLICANT: Zamudio, Carlos
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                   TYPE: PRT
1 PLSSIFSRIG 10
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98 PLTSIYNRLG 107
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                                                                       Local Similarity 60. hes 6; Conservative
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                                                                                       60.7%;
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                                                                                       Score 37;
Pred. No.
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Pred. No.
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RESULT 10 PCT-US02-32727-2247

Sequence 2247, Application PC/TUS0232727 GENERAL INFORMATION:

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TILE OF INVENTION: Compositions and Methods for FILE REFERENCE: 210121.514C1
CURRENT APPLICATION NUMBER: PCT/US02/32727
CURRENT FILING DATE: 2002-10-11
NUMBER OF SEQ ID NOS: 30992
SEQ ID NO 2247
LENGTH: 53
TYPE: PRT
ORGANITATION NUMBER OF SEQ ID NOS: 30992
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Best Local Similarity
Thas 7; Conserve
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PCT-US02-32727-2247
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US-10-057-498-2247
                                                                                                                                                                    RESULT 12
PCT-US02-32727-19192
                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Mitcham, Jennifer
APPLICANT: Mitcham, Jennifer
APPLICANT: Steiky, Yasir
APPLICANT: Persing, David
TITLE OF INVENTION: Compositions and Methods for the Therapy and Diagnosis of Acnes
FILE REFERENCE: 210121.514
CURRENT APPLICATION NUMBER: US/10/057,498
CURRENT FILING DATE: 2001-04-20
NUMBER OF SEQ ID NOS: 29212
SEQ ID NO 2247
LENGTH: 53
TYPE: PRT
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                                                   APPLICANT: Mitcham, Jennifer
APPLICANT: Skeiky, Yasir
APPLICANT: Skeiky, Pavid
APPLICANT: Persing, David
APPLICANT: Bhatis, Ajay
APPLICANT: Maisonneuve, Jean Francois
                                           APPLICANT:
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Skeiky, Yasir
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Jen, Shyian
Lodes, Michael
Wang, Sayian
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Bhatia, Ajay
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                        Zhang, Yanni
Wang, Siging
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70.0%;
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70.0%;
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; Pred. No. 18;
1; Mismatches
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Pred. No. 18;
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US-10-057-498-19192
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GENERAL INPORMATION:
GENERAL INPORMATION:
GENERAL INPORMATION:
GENERAL INPORMATION:
Miccham, Jennifer
APPLICANT: Miccham, Jennifer
APPLICANT: Persing, David
TITLE OF INVENTION: Compositions and Methods for the Therapy and Diagnosis of Acnes Vt.
FILE REFERENCE: 210121.514
CURRENT APPLICATION NUMBER: US/10/057,498
CURRENT FILING DATE: 2001-04-20
NUMBER OF SEQ ID NOS: 29212
SEQ ID NO 19192
LENGTH: 70
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LENGTH: 70
TYPE: PRT
ORGANISM: Propioni acnes
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Best Local
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APPLICANT: Jones, Robert
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
APPLICANT: Bouglass, John
TITLE OF INVENTION: Compositions and Methods for the Therapy and Diagnosis of Acnes
FILE REFERENCE: 210121.514C1
CURRENT APPLICATION NUMBER: PCT/US02/32727
CURRENT FILICATION NUMBER: PCT/US02/32727
                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 7; Conserv
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                                   APPLICANT
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T: Douglass, John
INVENTION: Compositions
RENCE: 210121.514C1
                                                                                Zhang, Yanni
Wang, Siqing
Jen, Shyian
Lodes, Michael
Benson, Darin
Jones, Robert
                                                                                                                                                                                     Maisonneuve, Jean
                                                  Carter, Darrick
Barth, Brenda
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Bhatia, Ajay
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77.8%;
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77.8%;
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Pred. No.
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Pred. No. 24;
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                 and Methods for the Therapy and Diagnosis of Acnes Vi
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RESULT 15
US-10-057-498-23079
US-10-057-498-23079
Sequence 23079, Application US/10057498
GENERAL INFORMATION:
APPLICANT: Mitcham, Jennifer
APPLICANT: Skeiky, Yasir
APPLICANT: Persing, David
TITLE OF INVENTION: Compositions and Methods for the Therapy and Diagnosis of Acnes V
FILLE REFERENCE: 210121.514
CURRENT APPLICATION NUMBER: US/10/057,498
CURRENT FILING DATE: 2001-04-20
UNMBER OF SEQ ID NOS: 29212
SEQ ID NO 23079
USCANISM: Propioni acnes
US-10-057-498-23079
Search completed: February 3, 2003, 09:35:01 Job time: 1.98258 secs
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CURRENT FILING DATE: 2002-10-11
NUMBER OF SEQ ID NOS: 30992
SEQ ID NO 23079
LENGTH: 144
TYPE: PAT
ORGANISM: Propioni acnes
PCT-US02-32727-23079
                                                                                                                                                       Query Match 57.4%; Score 35; DB 6; Length 144; Best Local Similarity 45.5%; Pred. No. 51; Matches 5; Conservative 4; Mismatches 2; Indels
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                                                                               2 LSSIFSRIGDP 12
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24 LAAVFKSLGDP 34
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Maximum DB
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Maximum Match 10
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Perfect score:
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          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 2000000000
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ALIGNMENTS

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C/Superfamily: hepatitis B virus surface antigen
C/Keywords: surface antigen
F/1-378/Product: surface antigen pre-S1 (large envelope protein) #status predicted <PS1>
F/1-108/Domain: pre-S1 domain #status predicted <PRE1>
F/109-378/Product: surface antigen pre-S2 (middle envelope protein) #status predicted <PEF)
F/109-378/Product: surface antigen pre-S2 (middle envelope protein) #status predicted <PEF)
F/109-163/Domain: pre-S2 domain #status predicted <PRE2>
F/164-378/Product: surface antigen S (small envelope protein) #status predicted <PSD>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            surface antigen - hepatitis B virus (subtype ayw, patient A) (fragment)
N;Alternate names: envelope protein, HBs antigen
N;Contains: surface antigen pre-S1 (large envelope protein); surface antigen pre-S2
C;Species: hepatitis B virus, HBV
A;Variety: subtype ayw, patient A
C;Date: 06-Jan-1995 #sequence_revision 01-Nov-1996 #text_change 26-Aug-1999
C:Date: 06-Jan-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Experimental source: subtype ayw, patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: EMBL:X77309; NID:g452611; PIDN:CAA54515.1; PID:g452612
                                                                                                                                                                                                                                                                                                                                                                                                                           A;Gene: S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-378 < LAI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Description: Sequence an
A; Reference number: S41869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 submitted to the EMBL Data Library, January 1994
A;Description: Sequence analysis of HBV genomes isolated from patients with HBsAg chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Accession: S41869
R;Lai, M.E.; Mazzole
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                                                     1 PLSSIFSRIGDP 12
PLSSIFSRIGDP 160
                                                                                                                                        Similarity
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                                                                                                                                  100.0%;
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Pred. No. 0.0011;
Nismatches 0;
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A; Molecule type: DNA A; Residues: 1-389 <GAL> A; Cross-references: GB:

GB:J02203

large surface antigen - hepatitis B virus (subtype ayw)
N;Contains: major surface antigen;
C;Species: hepatitis B virus, HBV
C;Date: 18-Dec-1981 #sequence revision 18-Dec-1981 #text_change 07-May-1999
C;Accession: A03703, JQ2064; FQOS91
R;Galibert, F.; Mandart, E.; Fitoussi, F.; Tiollais, P.; Charnay, P.
Nature 281, 646-650, 1979
A;Title: Nucleotide sequence of the hepatitis B virus genome (subtype ayw) i
A;Reference number: A93214; MUID:81012091; PMID:399327
A;Accession: A03703

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A;Gene: S
C;Superfamily: hepatitis B virus surface antigen
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: surface antigen
F;1-389/Product: surface antigen pre-S1 (large envelope protein) #status predicted <PSD>
F;164-389/Product: surface antigen S (small envelope protein) #status predicted <PSD>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A,Experimental source: genogroup D, subtype ayw3
R;Norder, H.; Courouce, A.M.; Magnius, L.O.
J. Gen. Virol. 73, 3141-3145, 1992
A;Title: Molecular basis of hepatitis B virus serotype variations within A;Reference number: PQ0453; MUID:93107848; PMID:1469353
A;Accession: PQ0591
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A; Residues: 164-389 < NOR >
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A;Title: Genetic relatedness of hepatitis B viral strains of diverse geographical origin A;Reference number: JQ2044; MUID:93329382; PMID:8336122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    submitted to the EMBL Data Library, March 1992
A;Description: Sequence analysis of HBV genomes isolated from patients wi A;Reference number: S20745
A;Recession: S20749
A;Molecule type: DNA
A;Residues: 1-389 <LAI>
A;Cross-references: EMBL:X65258; NID:g59434; PIDN:CAA46353.1; PID:g59435
A;Gross-references: EMBL:X65258; NID:g59434; PIDN:CAA46353.1; PID:g59435
A;Bxperimental source: subtype ayw, patient CI
A;Note: typical methionine for the beginning of pre-S2 domain at position C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Surface antigen - hepatitis B virus (subtype ayw, patient CI)
N;Alternate names: envelope protein; HBs antigen
N;Contains: surface antigen pre-Si (large envelope protein); surface antigen
C;Species: hepatitis B virus, HBV
A;Variety: subtype ayw, patient CI
C;Date: 20-Feb-1995 #sequence_revision 01-Nov-1996 #text_change 26-Aug-1999
C;Accession: S20749
B;Lai, M.E.; Mazzoleni, A.P.; Balestrieri, A.; Melis, A.; Porru, A.
submitted to the EMBL Data Library, March 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Keywords: glycoprotein; surface antigen (gene pre-S2/S) #status predicted F;109-389/Product: middle surface antigen (gene S) #status predicted F;164-389/Product: major surface antigen (gene S) #status predicted <MSA>F;164-389/Product: major surface antigen (gene S) #status predicted F;4,112,166/Binding site: carbohydrate (Asn) (covalent) #status predicted
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C;Superfamily: hepatitis B virus surface antigen
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A; Residues: 264-343 < NO2>
A; Experimental source: sul
C; Genetics:
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large surface antigen - hepatitis B virus (subtype ayw, N; Contains: major surface antigen; middle surface antige C; Species: hepatitis B virus, HBV
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Pred. No. 0.0012;
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Pred. No. 0.0012;
Mismatches 0
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                                                                                                              RESULT
SAVLA1
                             large surface antigen - hepatitis B virus (strain alphal) N;Contains: major surface antigen; middle surface antigen C;Species: hepatitis B virus, HBV
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A;Note: host Homo sapiens (man)
C;Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_change
C;Accession: A03704; PQ0585
R;Bichko, V.; Pushko, P.; Dreilina, D.; Pumpen, P.; Gren, E.
FEBS Lett. 185, 208-212, 1985
A;Title: Subtype ayw variant of hepatitis B virus: DNA primary statements number: A05237; MUID:85204397; PMID:3996597
A;Accession: A03704
A;Molecule type: DNA
A;Residues: 1-389 <BIC>
A;Residues: 1-389 <BIC>
A;Residues: 1-389 <BIC>
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A,Title: Molecular basis of hepatitis B virus serotype variations A;Reference number: PQ0453; MUID:93107848; PMID:1469353
A,Accession: PQ0585
A,Molecule type: DNA
A,Residues: 264-343 <NOR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                    large surface antigen - hepatitis B virus (subtype adyw)
N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBV
C;Species: hepatitis B virus, HBV
C;Accession: A93217; A03703
R;Pasek, M.; Goto, T.; Gilbert, W.; Zink, B.; Schaller, H.; MacKay, P.; Lead
Nature 282, 575-579, 1979
A;Title: Hepatitis B virus genes and their expression in B. coli.
A;Reference number: A93217; MUID:81012115; PMID:399329
A;Accession: A93217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Keywords: glycoprotein; surface antigen (gene pre-S2/S) #status predicted F;109-389/product: middle surface antigen (gene pre-S2/S) #status predicted c/MSA> F;164-389/product: major surface antigen (gene S) #status predicted c/MSA> F;4,112,166/Binding site: carbohydrate (Asn) (covalent) #status predicted
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SAVLAJ
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A;Experimental source: subtype
                                                                                                                                                                                                   A;Gene: pre-S1/pre-S2/S
C;Superfamily: hepatitis B virus surface antigen
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen
F;109-389/Product: middle surface antigen (gene pre-S2/S) #status predicted
F;164-389/Product: major surface antigen (gene S) #status predicted <MSA>
F;164-389/Product: major surface antigen (gene S) #status predicted <MSA>
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                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: DNA
A;Residues: 1-389 <PAS>
A;Cross-references: GB:J02202
C;Genetics:
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Best Local (
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149 PISSIFSRIGDP 160
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                                                  PLSSIFSRIGDP 12
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nilarity 91.7%;
Conservative 1
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        160
                                                                                                                                96.7%;
91.7%;
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Pred. No.
                                                                                                     1; Mismatches
                                                                                                                                Score 59; DB 1;
Pred. No. 0.0028;
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A;Gene: ...
A;Introns: 111/3
C;Superfamily: hepatitis B virus surface antigen
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: surface antigen pre-S1 (large envelope protein)
F;1-389/Product: surface antigen pre-S1 redicted <PRE1>
F;1-108/Domain: pre-S1 domain #status predicted envelope protein
F;109-389/Product: surface antigen pre-S2 (middle envelope protein)
F;109-389/Product: surface antigen pre-S2 (middle envelope protein)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Variety: Subtype ayw4
C;Date: 23-Nov-1994 #sequence_revision 12-May-1995
C;Accession: S47407; JQ2071
                                                                                                                                                                                                                                                                                      A; Molecule type: DNA
A; Residues: 164-389 < NOR>
                                                                                                                                                                                                                                                                                                                                                                                J. Gen. Virol. 74, 1341-1348, 1993
A;Title: Genetic relatedness of hepatitis B viral strains
A;Reference number: JQ2044; MUID:93329382; PMID:8336122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           surface antigen - hepatitis B virus (subtype ayw4)
N;Alternate names: envelope protein; HBs antigen
N;Contains: surface antigen pre-S1 (large envelope protein); surface antigen pre-S2
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A;Residues: 1-389 cTON>
A;Residues: 1-389 cTON>
A;Cross-references: EMBL:M32138; NID:g329667; PIDN:AAA45502.1; PID:g329669
A;Cross-references: EMBL:M32138; NID:g329667; PIDN:AAA45502.1; PID:g329669
R;Norder, H.; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I
J. Gen. Virol. 74, 1341-1348, 1993
J. Gen. Virol. 74, 1341-1348, 1993
A;Title: Genetic relatedness of hepatitis B viral strains of diverse geogra
A;Reference number: JQ2044; MUID:93329382; PMID:8336122
A;Accession: JQ2082
A;Accession: JQ2082
                                                                                                                                                                                                                                           C;Genetics:
                                                                                                                                                                                                                                                                                                                                           A; Accession: JQ2071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 1-389 <PLU>
A;Cross-references: EMBL:Z35716; NID:g527435; PIDN:CAA84788.1;
A;Experimental source: subtype ayw4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         submitted to the EMBL Data Library, A;Description: Molecular cloning and
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A;Title: Active hepatitis B virus replication in the presence A;Reference number: A34773; MUID:90266476; PMID:2345966
A;Accession: B34773
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 164-389 < NOR>
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C;Accession: B34773; JO2082
                                                                                                                                                                                                                                                                                                                                                                                                                                        A;EXPERIMENTAL BOURCE: SUBLYPE AYWA

1;EXPERIMENTAL BOURCE: SUBLYPE AYWA

1;NORDER, H.; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I.K.; Magnius

1, Gen. Virol. 74, 1341-1348, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Molecule type:
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;109-389/Product: middle surface antigen (gene pre-S2/S) #status predicted
;164-389/Product: major surface antigen (gene S) #status predicted <MSA>
;4,112,166/Binding site: carbohydrate (Asn) (covalent) #status predicted
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Similarity 91.7%;
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Pred. No. 0.
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                                  envelope protein) #status predicted
<PRE2>
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                       Query Match
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Matches 10
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A;Introns: 123/2
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen
                                                                                                                                                                                                                                                                                                                               N;Alternate names: envelope protein
N;Contains: major surface antigen;
C;Species: hepatitis B virus, HBV
A;Variety: isolate 29Y11HCC
                                                                                  A; Experimental C; Genetics:
                                                                                                                                                                                                Arch. Virol. 143, 2313-2326, 1998
A;Title: Hepatitis B virus genomic sequence in the circulation A;Reference number: Z17684; MUID:99129050; PMID:9930189
A;Accession: T13474
                                                                                                                                                                                                                                                        C;Date: 13-Aug-1999 #sequence_revision 13-Aug-1999 #text_change C;Accession: T13474 R;Takahashi, K.; Akahane, Y.; Hino, K.; Ohta, Y.; Mishiro, S. Arch. Virol. 143, 2313-2326, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Reywords: surface antigen
C;Reywords: surface antigen
F;1-389/Product: large surface antigen (gene pre-S1/pre-S2/S) #status predicted <DSL>
F;109-389/Product: middle surface antigen (gene pre-S2/S) #status predicted <DSA>
F;164-389/Product: major surface antigen (gene S) #status predicted <MSA>
                                                                                                                        A;Residues: 1-382 <TAK>
A;Cross-references: EMB
                                                                                                                                                         A; Molecule type: DNA
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A:Variety: subtype ayw, isolate patient C1005
C:Date: 20-Feb-1995 #sequence_revision 06-Dec-1996 #text_change 06-Dec-1996
C:Accession: S32202
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S32202
                                                                  A;Gene: S
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A;Cross-references: EMBL:X72702
A;Experimental source: subtype
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submitted to the EMBL
A:Description: Identif
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        large surface
;Superfamily: hepatitis I
;Keywords: glycoprotein;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
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                                                                                                                                                                          preliminary; translated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PISSIFSRIGDP 160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antigen - hepatitis B virus
ajor surface antigen; middle
patitis B virus, HBV
                                                                                                     source:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ne EMBL Data Library
Identification and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S.; Schlayer, M.J.; Peters, T.;
                                                                                                                                                                                                                                                                                                                                                                             envelope protein; HBs antigen
surface antigen; middle surface
                                                                                           EMBL:AB014388; NID:g3582381; PIDN:BAA32936.1; PID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12
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91.7%;
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91.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 59;
Pred. No.
                                                                                                                                                                            from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 59; DB 2;
Pred. No. 0.0028;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence analysis of hepatitis B virus DNA in immunolog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           March
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                                                                                                                                                                                                                                                                                                                                                                               surface
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2;
0.0028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           envelope protein) #status predicted <SAG:
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                                                                                                             PID: 93582384
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                                                                                             29Y11HCC
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10;

Conservative

Similarity

88.5**%**; 83.3**%**;

; Score 54; DB; Pred. No. 0.02

8

Length 382;

9

Gaps

0

DB 2;

surface

antigen

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A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-384 <TAK>
A;Residues: 1-384 <TAK>
A;Cross-references: EMBL-AB014368; NID:g3551304; PIDN:BAA32864.1; PID:g3551308
A;Experimental source: Japanese patient with hepatocellular carcinoma isolate 09D09HCC
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N, Contains: major surface antigen;
C, Species: hepatitis B virus, HBV
A, Variety: isolate 09D09HCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Date: 13-Aug-1999 #sequence_revision 13-Aug-1999 #text_change 21-Jul-2000 C;Accession: T13469
R;Takahashi, K.; Akahane, Y.; Hino, K.; Ohta, Y.; Mishiro, S. Arch. Virol. 143, 2313-2326, 1998
A;Title: Hepatitis B virus genomic sequence in the circulation of hepatocell A;Title: Hepatitis B virus genomic sequence in the circulation of hepatocell A;Reference number: Z17684; MUID:99129050; PMID:9930189
A;Accession: T13469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nicontains: surface antigen pre-Si (large envelope protein); surface antiger c; Species: hepatitis B virus, HBV A; Variety: subtype ayw, patient C C; Date: 20-Feb-1955 #sequence_revision 01-Nov-1996 #text_change 26-Aug-1999 C; Accession: $20745 R; Lai, M.E.; Mazzoleni, A.P.; Balestrieri, A.; Melis, A.; Porru, A. submitted to the EMBL Data Library, March 1992 A, Description: Sequence analysis of HBV genomes isolated from patients with A; Reference number: $20745
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Introns: 123/2
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             large surface antigen - hepatitis B virus (isolate 09D09HCC:
N,Alternate names: envelope protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              문
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                                                                                                                                                                                                                                                         A;Molecule type: DNA
A;Residues: 1-389 <LAAI>
A;Cross-references: EMBL:X65257; NID:g59429; PIDN:CAA46349.1; PID:g59430
A;Cross-references: EMBL:X65257; NID:g59429; PIDN:CAAA6349.1; PID:g59430
A;Cross-references: EMBL:X65257; NID:g59429; PID:g59430
A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  surface antigen - hepatitis B virus (subtype ayw, patient
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                                                                                                                                               Local Similarity
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2 LSSIFSRIGDP
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                                                                                                              Conservative
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Pred. No. 0.02
1; Mismatches
                                                                                                              0
                                                                                                                                                        Score 54;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    middle surface antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (large envelope protein); surface antigen
                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genomes isolated from patients with HBsAg negat
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                                                                                                                                                        0.025;
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                                                                                                                                                                                          Length 389
                                                                                                                  Indels
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large surface antigen - hepatitis B virus (subtype adr)
N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBV
C;Species: hepatitis B virus, HBV
C;Accession: A03705; S04569; JQ2107; PQ0608
C;Accession: A03705; S04569; JQ2107; PQ0608
R;Ono, Y.; Onda, H.; Sasada, R.; Igarashi, K.; Sugino, Y.; Nishioka, K.
Nucleic Acids Res. 11, 1747-1757, 1983
A;Title: The complete nucleotide sequences of the cloned hepatitis B virus IA;Reference number: A93460; MUID:83168919; PMID:6300776
A;Accession: A03705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: GB:V00867

A;Cross-references: GB:V00867

A;Rho, H.M.; Kim, K.; Hyun, S.W.; Kim, Y.S.

Nucleic Acids Res. 17, 2124, 1989

A;Title: The nucleotide sequence and reading frames of a mutant hepatitis B virus subtype
A;Reference number: S04568; MUID:89183619; PMID:2928116

A;Accession: S04569
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A;Title: Genetic relatedness of hepatitis B viral strains of diverse geographica A;Reference number: JQ2044; MUID:93329382; PMID:8336122
A;Accession: JQ2107
A;Accession: JQ2107
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R,Norder, H.; Courouce, A.M.; Magnius, L.O.
J. Gen. Virol. 73, 3141-3145, 1992
J. Gen. Virol. 73, 3141-3145, 1992
A,Title: Molecular basis of hepatitis B virus serotype variations within the four major :
A,Reference number: PQ0453; MUID:93107848; PMID:1469353
                                                                                                                                                                                                                                                    A;Gene: pre-S1/pre-S2/S
A;Gene: pre-S1/pre-S2/S
C;Superfamily: hepatitis B virus surface antigen
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen (gene pre-S2/S) #status predicted
F;120-400/Product: middle surface antigen (gene S) #status predicted <MSA>
F;175-400/Product: major surface antigen (gene S) #status predicted <MSA>
F;15,123,177/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Molecule type: DNA
A;Residues: 1-50,'R',52-66,'YP',69-129,'V',131-142,'P',144-164,'S',166-176,'S',178-338,'C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A, Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                               A/Experimental source: subtype adrq+, C/Genetics:
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A; Residues: 175-400 < NOR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 275-354 < NO2>
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                                                                                                                               Matches
                                                                                                                                                             Query Match
Best Local
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160
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                                                            PLSSIFSRIGDP 12
PISSIFSRTGDP
                                                                                                                               l Similarity
10; Conservat
                                                                                                                               Conservative
                                                                                                                                                             83.3%;
                                                                                                                               1; Mismatches
                                                                                                                                                                Score 54; DB 1;
Pred. No. 0.026;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bau
                                                                                                                                                                                              Length 400;
                                                                                                                                         Indels
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                                                                                                                                         0
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N;Alternate names: envelope protein; HBs antigen N;Contains: surface antigen pre-S1 (large envelope C;Species: hepatitis B virus, HBV A;Variety: subtype adr C;Date: 09-Dec-1993 #sequence_revision 10-Nov-1995 C;Accession: S35528

surface antigen - hepatitis B virus (subtype adr) N;Alternate names: envelope protein; HBs antigen

R;Mukaide, M.; Kumazawa, T.; Hoshi, A.; Kawaguchi, R.; Hikiji, Nucleic Acids Res. 20, 6105, 1992

e E

hepatitis B

virus,

aubtype

adr

(SRADR)

ج

#text_change

protein); surface antigen

pre-S2

(mide

The complete nucleotide

prm)

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A/Introns: 210/3
C/Superfamily: hepatitis B virus surface antigen
C/Keywords: surface antigen
C/Keywords: surface antigen
P/1-445/Product: surface antigen pre-S1 (large envelope protein) #status
                                                                                                                                                                                                                                                                                                                                                                                                A;Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Loncarevic, I.F.; Zentgraf, H.; Schroeder, C.H.
Nucleic Acids Res. 18, 4940, 1990
A;Tille: Sequence of a replication competent hepatitis B
A;Reference number: S12598; MUID:90370503; PMID:2395664
A;Accession: S43492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      surface antigen - hepatitis B virus (subtype adr)
N;Alternate names: HBs antigen
N;Contains: surface antigen pre-S1 (large envelope protein); surface antigen pre-S2 (mi
C;Species: hepatitis B virus, HBV
A;Variety: subtype adr
C;Date: 07-Sep-1994 #sequence_revision 10-Nov-1995 #text change 26-Aug-1999
                                                                                                                                                                                                                                                                                                                                                 A;Gene:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Mimms, L.T.; Solomon, L.R.; Ebert, J.W.; Fields, H.
Biochem. Biophys. Res. Commun. 195, 186-191, 1993
A;Title: Unique PreS sequence in a gibbon-derived hepatitis B virus variant.
A;Reference number: PN0601; MUID:93371402; PMID:8363598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
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A;Experimental source: subtype adr
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, August 1992
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-400 < M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Gene: S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Reference number:
A;Accession: S35528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Experimental source: subtype adr.
Mimms, L.T.; Solomon, L.R.; Ebert,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Introns: 165/2
                                                                                                                                         1-164/Domain: pre-S1 domain #status predicted <PREI>
;165-445/Product: surface antigen pre-S2 (middle envelope protein)
;165-219/Domain: pre-S2 domain #status predicted <PRE2>
;20-445/Product: surface antigen S (small envelope protein) #status
                                                                                                                                                                                                                                                                                                                                                                                              ;Molecule type: protein ;Residues: 165-196 <MI2>
                                                                           Query Match
Best Local |
                                                                                                                                                                                                                                                                                                                                                                                                                                               Accession: PN0602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;Residues: I-445 <LON>
;Cross-references: EMBL:X52939; NID:g457780; PIDN:CAA37114.1; PID:g457783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ',1-400/Product: surface antigen pre-S1 (large envelope protein) #status predicted <PS;1-119/Domain: pre-S1 domain #status predicted <PRE1>;120-400/Product: surface antigen pre-S2 (middle envelope protein) #status predicted ',120-174/Domain: pre-S2 domain #status predicted <PRE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Date: 07-Sep-1994 #sequence revision 10-Nov-1995 #text_change 26-Aug-1999
Accession: S43492; PN0601; PN0602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Status: translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           175-400/Product: surface antigen S (small envelope protein) #status predicted <PSD>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Superfamily:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          160 PISSIFSRIGDP 171
1 PLSSIFSRIGDP 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10;
                                                                        Similarity
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                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <MIM>
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                                                                88.5%;
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                                                                   Score 54; I
Pred. No. 0.
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Pred. No.
                                            Mismatches
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                                                                DB 2;
0.029;
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Search completed: February Job time: 1.79443 secs

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2003, 09:26:29

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C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen
F;1-389/Product: surface antigen pre-S1 (large envelope protein) #status predicted <PS1>
F;1-108/Domain: pre-S1-specific domain #status predicted <PRE1>
F;109-389/Product: surface antigen pre-S2 (middle envelope protein) #status predicted <P:
F;109-163/Domain: pre-S2-specific domain #status predicted <PRE2>
F;164-389/Product: surface antigen S (small envelope protein) #status predicted <PSD>
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A;Title: Complete sequencing of a gibbon hepatitis B virus genome reveals a unique genoty A;Reference number: $67503; MUID:96207410; PMID:8615024
A;Accession: $67506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: hepatitis B vīrus, HBV
A;Variety: subtype ayw3, isolate Hope CH1357
C;Date: 29-Jan-1998 #sequence_revision 06-Peb-1998
                                                                                                                                                                                                   F;4,112,166/Binding site:
                                                                                                                                                                                                                                                                                                                                                                                                          A;Gene: pre-S1/pre-S2/S
A;Introns: 112/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: EMBL:U46935; NID:g1814218; PIDN:AAB41952.1; A;Experimental source: subtype ayw3, isolate Hope CH1357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: DNA
A; Residues: 1-389 < NOR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Accession: S67506
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       large surface antigen - hepatitis B virus (subtype ayw3,
                                                                                                                         Query Match
Best Local :
                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genetics:
150 ISSIPSRTGDP 160
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                                               2 LSSIFSRIGDP
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                                                                                                                            Similarity
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                                                                                                                    77.0%;
81.8%;
                                                                                                                                                                                                 carbohydrate (Asn)
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                                                                                            Score 47; DB 2;
Pred. No. 0.54;
1; Mismatches
                                                                                                                                                                                              (covalent) #status
                                                                                                                                            Length 389;
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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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A Geneseq 101002:*

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Gapop 10.0 , Gapext 0.5
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831.368 Million cell updates/sec
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Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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174	174	164	164	102	60	36	25	12	12	Length	
13	12	21	21	20	16	23	12	21	21	B	
AAR23867	AAR15617	AAY90760	AAY93740	AAY30947	AAR76747	AAE17023	AAR14747	AAY87906	AAB10598	ij	SUMMARIES
Pre-S gene region	HBBAG pre-S region	Henatitis B virus	Amino acid semienc	Himan Hhead (Drace	HRV surface antice	Hepatitis B virus	HBV Pre S2 pentide	HBV cell surface n	HBV fusion protein	Description	

RRXFF

WPI; 2000-514959/46. N-PSDB; AAA71751.

Particle for cell-specific gene delivery, useful in gene therapy, comprises nucleic acid in protein coat that includes a fusion protein

45	4	43	42	4.	40	. 39	. u	37	36	35	34	w	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	Ħ
51	52	52	52	52	52	52	54	5.4	54	54	54	54	54	54	54	51	54	54	54	59	59	59	59	59	59	59	59	59	13	61	61	13	61	61
			•				88.5			88.5			_								96.7	96.7	96.7	96.7	96.7	•	•	96.7	100.0	100.0	100.0	•	100.0	
175	399	399	389	174	55	54	400	400	389	281	281	281	281	281	281	281	174	174	174	389	347	174	174	67	67	64	6 <u>4</u>	55	393	389	389	281	281	215
10	22	22	2	10	22	15	21	17	7	22	80	15	œ	7	7	7	ü	12	12	22	21	13	12	23	22	23	22	œ	22	IJ	13	20	13	21
AAP91482	AAG66932	AAG66919	AAG62931	AAP91483	AAE04710	AAR47945	AAY54045	AAR93801	AAP60794	AAM48419	AAY32834	AAR62870	AAP70294	AAP60617	AAP60560	AAP60163	AAR23871	AAR15622	AAR12394	AAG66931	AAB10596	AAR23868	AAR15618	AAE17029	AAB81135	AAE17027	AAB81133	AAP70002	AAU14097	AAR27474	AAR27472	AAY32835	AAR27471	AAB10597
nslation pro	genoty	7		9 (vi ma	oinant	20.	-	henatiti) (ч -	٦. ر م	B William	m .	gene r	Ad pre-S regi	-Santige	genoty	<u> </u>	Cone t	HBRAG DYA-S TATION	minally	D (Henatitis B virus	p Š		titia R virus	700000000000000000000000000000000000000	protein		Ag protein	HBV fusion protein

ALIGNMENTS

RESULT 1	년 1
Ę	AAB10598 standard; Protein; 12 AA.
	AAB10598;
	08-JAN-2001 (first entry)
DE I	HBV fusion protein cell permeability peptide fragment.
	N 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
	Fusion protein; protein coat; virus-specific packaging signal, aci.
	virus protein; cell permeability; cell-specific binding signal; ps;
₹ ₹	large surface protein; core antigen; gene therapy.
	Hepatitis b virus
	WO200046376-A2.
	10-AUG-2000.
	04-F6B-2000; 2000W0-DE00363.
	05~FEB-1999; 99DE-1004800.
	(HILD/) HILDT B.
PI H	Hildt E, Hofschneider P:
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of viral

protein,

permeability peptide and cell-binding site

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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel particle (A), comprising a protein coat with a fusion protein (FP), and, inside the coat, a nucleic acid (I) including the sequence for a virus-specific packaging signal (psi) and a structural gene. FP contains a virus protein (VP), a peptide (P) that mediates cell permeability and a heterologous cell-specific binding site (RGD). The invention also describes (1) producing (A) in which FP contains an LHBs (large surface protein of hepatitis B virus (HBCN)) and RGD; (2) preparing (A) in which FP contains an HBV core antigen (HBCAG), (P) and RGD; (3) FP; (4) DNA encoding FP; and (5) expression vector containing the DNA of (d). The products of the invention are used in gene therapy of cells and tissues, in vivo or ex vivo. This sequence represents a fragment of the Hepatitis b virus cell permeability peptide which is described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HBV cell surface protein ZPP fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY87906;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY87906 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cell surface protein;
This invention describes a novel cell permeability mediating polypeptide (CPM) comprising (A) or a sequence which differs by one or more amino acids, which shows cell permeability and is not hepatitis B virus (HBV) surface protein and where the DNA sequence of the latter amino acid sequence hybridizes with the DNA of (I). CPM can be used to mediate cell permeability of substances, especially polypeptides, nucleic acids and chemicall connections. This is useful for increasing the reactivity of such substances in theraspeutic treatment of various conditions. This sequence represents a fragment of the HBV cell surface protein ZPP which
                                                                                                                                                                                                                                                                                                                                        03-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                      03-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                  18-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                DE19850718-C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy; treatment.
                                                                                                                                                                                  Novel cell permeability-mediating polypeptide useful permeability of substances such as other polypeptides
                                                                                                                                                                                                                               N-PSDB, AAA39621
                                                                                                                                                                                                                                                WPI; 2000-340689/30
                                                                                                                                                                                                                                                                                                           (HILD/) HILDT
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milarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                                                         German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZPP; cell permeability mediating polypeptide;
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Pred. No. 3.7e-05;
, Mismatches 0;
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RXRXRX

(first entry)

AAE17023 standard; peptide; 36 AA

Hepatitis B 18-APR-2002 AAE17023;

(HBV) ayw

subtype pre-S2

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ARSULT 3
AAX1477
ID AAX147
XX AAX1
XX AAX1
XX MONC
KW 
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RESULT 4
AAE17023
ID AAE1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           epitope; antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monoclonal antibodies; hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HBV Pre S2 peptide
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                                                                                                                                                                                                                                                                                                                                                                           The peptide corresponds to residues 150-174 of the HBV PreS2 protein. It was used to raise monoclonal antibodies specific for the HBV M protein but which do not bind to the HBV S or L proteins. The MAD can be used to sub-type HBV and to prepare subunit voaccines for HBV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-NOV-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New MADs to PreS2 and PreS1 polypeptide(s) of Hepatitis-Benyelope - used to sub-type HBV in sample and as vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1991-334198/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mimms LT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAY-1990;
                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                             11 PLSSIFSRIGDP
                                                                                                                                                                            1 PLSSIFSRIGDP 12
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12; Conser
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12; Conser
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                                                                                                                                                                                                                         100.0%; milarity 100.0%; loonservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                    AA.
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Pred. No. 3.7e-05;
                                                                                                                                                                                                                                                           Score 61; DB 12;
Pred. No. 8.3e-05;
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RESULT 5
AAR76747
ID AAR7
XX
AC AAR7
AC AAR7
AC 18-M
DT 18-M
XX
DB HBV
XX
                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                          carrier to present the epitope to the immune system. This chimeric probhylactic or therapeutic vaccination of the human or animal body, particularly against HBV. The nucleic acid encoding the protein may be used in gene therapy or DNA vaccination protocols. The chimeric probhylactic vaccine against a range of disease, e.g. HBV, hepatitis A virus (HAV), hepatitis C virus (HCV), influenza, foot-and-mouth disease, polio, herpes, rabies, acquired immunodeficiency syndrome (ALDS), dengue fever, malbers, acquired immunodeficiency syndrome (ALDS), salmonellosis, typhoid, food poisoning, diarrhoea, meningitis or gonorrhea. The present sequence is Hepatitis B virus (HBV) ayw
      HBV surface antigen pre-S2 region and FimH residues 224-226.
                                               18-MAR-1996
                                                                               AAR76747
                                                                                                          AAR76747 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to modified proteins comprising hepatitis B virus (HBV) core antigen (HBcAg) wherein one or more of the four arginine repeats has been deleted and the protein comprising the C-terminal cysteine of HBcAg. The deleted region may be replaced by an epitope from a protein other than HBcAg, in which case the HBcAg at a sia
                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                       subtype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 3A; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-098223/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-JUN-2000; 2000GB-0015308, 06-OCT-2000; 2000GB-0024544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis B virus; HBV; core antigen; HBcAg; immune system; typhoid; prophylactic; gene therapy; vaccine; hepatitis A virus; HAV; herpes; hepatitis C virus; HCV; influenza; foot-and-mouth disease; diarrhoea; tuberculosis; polio; rabies; acquired immunodeficiency syndrome; AIDS; dengue fever; yellow fever; malaria; whooping cough; salmonellosis; food poisoning; meningitis; gonorrhea; antiviral; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-JUN-2001; 2001WO-GB02817
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                                                                                                                                                                                        1 PLSSIFSRIGDP 12
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22 PLSSIFSRIGDP 33
                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   vaccine
                                                                                                                                                                                                                                                           12;
                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                     pre-S2 region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ins comprising a modified hepatitis B core antigen, useful as in prophylactic or therapeutic vaccination of the human or dy, particularly against hepatitis B virus infection -
                                                                                                                                                                                                                                                                                                                        36 AA;
                                                                                                                                                                                                                                                        Conservative
                                             (first entry)
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                                                                                                                                                                                                                                               Score 61; DB 23,
Pred. No. 0.00012;
                                                                                                                                                                                                                                                                                Length 36;
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AAY30947;

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RESULT 6
AAY30947
ID AAY3
XX
AC AAY3
XX
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                                                                                                                                                                                                                                        CC This sequence is encoded by a fragment of the plasmid pLPA38, which CC comprises a finH fusion gene comprising the sequence encoding the pre-S2 CC region of the hepatitis B viral surface antigen inserted into the finH CC gene. The primer sequences given in AAQ93057-58 caused the insertion of CC the pre-S2 region into the FinH protein at position 224-226. The CC resulting chimeric gene was then further modified by insertion of the CC type 1 fimbriae. Restiction sit the free position of the FinH adhesin of CC type 1 fimbriae. Restiction sit he pitopes are then inserted in-frame. CC into the finH gene, and the foreign epitopes are then inserted in-frame. CC in the selected positions the insertion of the epitopes did not captesion of the chimeric proteins on the surface of fimbriae on CC significantly alter the adhesive function of the FinH protein. The CC expression of the chimeric proteins on the surface of fimbriae on CC bacterial hosts illustrated the possibility of using bacterial adhesins companies and proteins which may be used in the production of variant FinH adhesins which may CC be useful for targetting active compounds and microbial cells to
                                                                                                                                                                              Matches
                                                                                                                                                                                               Query Match
Best Local :
                                          AAY30947 standard;
                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence is encoded by a fragment of the plasmid pLPA38,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Page 57; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Receptor specific bacterial adhesins - useful compounds and microbial cells to locations of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; AAQ93059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hasty DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-JAN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FimH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       W09520657-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GXBI-) GX BIOSYSTEMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chimeric -
                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                     42
                                                                                                                                              1 PLSSIFSRIGDP 12
                                                                                                                   PLSSIFSRIGDP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1995-275442/36.
                                                                                                                                                                              12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       type 1 fimbriae; organelle; adhesin; alpha-D-mannoside residue; FimF; FimG; receptor binding site; PCR; amplify.
                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                      60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Klemm P, Molin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis B virus.
Escherichia coli.
                                                                                                                                                                       100.0%; Score 61; DB 16; Length 60; Larity 100.0%; Pred. No. 0.00021; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94US-0187166.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note=
                                                                                                                 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'note=
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                                         Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Represents FimH residue 226"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Linker peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "HBV surface antigen pre-S2 region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Linker peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Represents FimH residue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sokurenko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for targetting active receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EV;
                                                                                                                                                                       0,
                                                                                                                                                                       Gaps
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RESULT 7
AAY93740
ID AAY9
                                                                                                                                                                                                                                                                                                                                                                         cc that modulates the immune response to a genetic vaccine vector (A), or cencodes a polypeptide (II) with similar effect. The invention also cc describes the identification of a polymclectide (Ia) encoding an cc accessory molecule (Ia) that improves transport and presentation of antigen by a cell. The products of the invention have antibacterial, cc antigen by a cell. The products of the invention have antibacterial, cc antivaral, antifungal, anti-altergic, antidiabetic, anti-inflammatory, canti-arthritic, anti-asthma, anticancer and immunomodulatory. Optimized (I) are incorporated into (A), or (I) or its encoded (II) are conditions (bacterial, viral or fungal), autoimmune disease (e.g. conditions (bacterial, viral or fungal), autoimmune disease (e.g. cc administered together with (A). (A) are used to treat or prevent cc athing or cancer (including metastases), inmune deficiency, allergy, cc sathma or cancer (including metastases), (I) are also used for cesponse to particular requirements, e.g. to direct a Thi-type helper cresponse to increase humoral or cellular responses (functioning as cc adjuvant) to control B or T cell proliferation to induce immunogolobulin control or isotype switching. This sequence represents the HBsAg creation to represents the HBsAg creation creation creation creation creation creation creation creation 
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammatory condition; psoriasis; immune deficiency; metastases; cancer; Thitype helper response; humoral; cellular response; B cell T cell; proliferation; immunogolobulin synthesis; isotype switching; immunomodulator; immune response; asthma; human; HbsAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 70; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying nucleic acid that directly or indirectly modulates the immune response to a genetic vaccine vector, e.g. for prevention of infection or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-508645/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Howard R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-FEB-1998;
11-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MAXY-) MAXYGEN INC
  AAY93740 standard; Protein; 164 AA
                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                           y Match 100.0%;
Local Similarity 100.0%;
hes 12; Conservative 0
                                                                                                                        4
                                                                                                                                                                     μ
                                                                                                                                                                                                                                                                                                                                                                  invention.
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                                                                                                                        PLSSIFSRIGDP
                                                                                                                                                                     PLSSIFSRIGDP 12
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                                                                                                                                                                                                                                                                                                                    102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             describes the identification of a novel polynucleotide (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-0074294.
98US-0021769.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stemmer
                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                            Score 61; DB 20;
Pred. No. 0.00038;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPC,
                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RG
                                                                                                                                                                                                                                                                     Length 102;
                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                            Gape
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Query Match
Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY93740;
                                                                                                                                                                                                                                                                                                                                                        The specification describes a plant expression vector which comprises two expression cassettes, the first comprising a polynucleotide encoding an antigen and the second comprising a non-identical polynucleotide encoding the same antigen. The antigen is especially a heparitis B surface antigen (HBBAG). The expression vector is used to transform bacterial and plant cells to elicit the production of anti-hepatitis B antibodies and are useful as vaccines. The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New expression vector for transforming plants comprising two expression cassettes useful for producing plant material comprising anti-hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-JUN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amino acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; AAA47000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mason
                                                                                                                                                                                                                                                                                                                                              a pre-S peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HEAL-) HEALTH RES
                                                           fusion protein; diagnosis; detection; variant; vaccine; chape coprecipitate.
                                                                                                                                                                              AAY90760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BOYC-) BOYCE THOMPSON INST PLANT RES
                                                                                                                                 18-AUG-2000
                                                                                                           Hepatitis B virus preS protein sequence
                                                                                                                                                                                                                                    149
                     WO200020606-A1
                                          Hepatitis B virus
                                                                                                                                                                                                                                                         .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000-452181/39.
                                                                                                                                                                                                                                     PLSSIFSRIGDP 160
                                                                                                                                                                                                                                                         PLSSIFSRIGDP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ,SH
                                                                                                                                                                              standard; Protein; 164 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thanavala Y,
                                                                                                                                                                                                                                                                                                                            164 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fig 5; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-0113827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US31020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for pre-S peptide
                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arntzen CJ,
                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                           Score 61; DB 21;
Pred. No. 0.00064;
                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Richter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E
                                                                                                                                                                                                                                                                                                      Length 164;
                                                                                                                                                                                                                                                                                    Indels
                                                                              chaperone;
                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                    Gaps
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02-OCT-1998;

98WO-EP06298

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RESULT 9
AAR15617
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
   Hepatitis B virus surface antigen epitope(s) immunogens or diagnostic reagents
                                        WPI; 1991-369007/50
                                                               Milich
                                                                                                          11-MAY-1990;
                                                                                                                               10-MAY-1991;
                                                                                                                                                                                                                      T-cell epitope;
                                                                                                                                                      28-NOV-1991.
                                                                                                                                                                             WO9117768-A.
                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                          HBsAg pre-S region subtype ayw.
                                                                                                                                                                                                                                                                                                             AAR15617 standard; protein; 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a polynucleotide (I) encoding a fusion protein (FP) (II) which is stable in a cell, and comprises a first polypeptide (P1) and a second polypeptide (P2) which co-precipitates a chaperone. The invention also provides methods for the production of fusion proteins which comprise epitopes. Compositions comprising the fusion proteins or epitopes of the invention are used as vaccines, and and efficient means of reliably expressing (poly) peptides comprising proteins. The present sequence represents the Hepatitis B virus (HBV) present sequence, which is used in an example from the present
                                                                                      (SCRI)
                                                                                                                                                                                                                                                                     25-MAR-1992
                                                                                                                                                                                                                                                                                            AAR15617;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                          150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 26; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel polynucleotide encod cell, useful for the produ chaperones, by using trunc with an intact N-terminus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-328936/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reimann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                             1 PLSSIFSRIGDP 12
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                                                               뮸
                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
12; Conserv
                                                                                  SCRIPPS CLINIC & RE
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SCHIRMBECK R.
                                                              Thornton
                                                                                                                                                                                                                                                                                                                                                                                                                                                               164 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schirmbeck R;
                                                                                                                                                                                                                                                                  (first entry)
                                                                                                         90US-0522663
                                                                                                                               91WO-US03268
                                                                                                                                                                                                                   vaccine; hepatitis B virus; antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-EP06298
                                                                                                                                                                                                                                                                                                                                                                       161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoding a fusion protein which is stable in a production of peptides which coprecipitate truncated variants of the SV40 large T antigen
                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                               Score 61; DB
Pred. No. 0.0
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                           0.00064;
                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 21;

    useful

                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 164;
                                                                                                                                                                                                                                                                                                                                                                                                                Indels
              as vaccines
                                                                                                                                                                                                                                                                                                                                                                                                              0;
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RESULT 10
ARZ3367
AZ AARZ3
AX AARZ3
XX Hepad
XX O'ALY
PM ('CALY
PM (NYBL
XX Hepad
XX
The sequences given in AAR23867 - AAR23871 are amino acid sequences C deduced from sequences of the pre-S portion of the env genes C corresponding to several hepatitis B virus (HBV) subtypes. These C proteins have properties distinct from those of the S-protein in C residues, an abscence of Cys residues, the highest subtype-dependant C variability among HBV DNA gene products and little homology with C analogous sequences corresponding to nonhuman hepadnavirus. These C properties suggest that the pre-S gene coded portion of the HBV C envelope is expose to the surface of the virion, is a target for the host immune response and is responsible for the host range of HBV C (limited to humans and some primates). Synthetic paptides and C antibodies against them, having predetermined specificity offer the C opportunity to explore the biological role of the pre-S protein moiety
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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Best Local s
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis B pre-S peptide immunogen and vaccine - for treatment and diagnosis of hepatitis {\tt B}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig 2; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-161100/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-APR-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-APR-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EP485361-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CALY ) CALIFORNIA INST TECHNOLOGY (NYBL-) NY BLOOD CENT INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitits B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            env gene; hepatitis B virus; HBV; S-protein; hydrophilic; hepadnavirus; immune response; antibody; envelope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR23867 standard; protein; 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pre-S gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR23867;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   also AAR15618-R15622.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              160 PLSSIFSRIGDP 171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 PLSSIFSRIGDP
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12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Neurath AR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       region translation product (1).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          virus, subtype ayw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 174;
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RESULT 11
AAB10597
ID AAB10
XX AAB10
XX AAB10
XX OB-JF
DT OB-JF
DX HBV I
XX Fusic
KW Virus
KW
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Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fusion protein; protein coat; virus-specific packaging signal; psi; virus protein; cell permeability; cell-specific binding site; LHB; large surface protein; core antigen; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB10597 standard; Protein; 215 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HBV fusion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JAN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis b virus.
                                                                                                                                                                                  with a fusion protein (FP), and, inside the coat, a nucleic acid (I) including the sequence for a virus-specific packaging signal (psi) and a structural gene. FP contains a virus protein (VP), a peptide (P) that mediates cell permeability and a heterologous cell-specific binding site (RCD). The invention also describes (I) producing (A) in which FP (RCD). The invention also describes (I) producing (A) in which FP (RCD). The propering (A) in which FP contains an LHBs (large surface protein of hepatitis B virus (HBV)) and (CONTAINS AND PR) (A) DNA encoding FP; and (5) expression vector containing the DNA of (d). The products of the invention are used in gene therapy of cells and tissues, in vivo or ex vivo. This sequence represents a fusion protein which is described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200046376-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-FEB-2000; 2000WO-DE00363.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprises nucleic acid in protein coat of viral protein, permeability peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Particle for cell-specific gene delivery, useful in gene therapy, comprises nucleic acid in protein coat that includes a fusion protein of viral protein, permeability peptide and cell-binding site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 14; Fig 2; 34pp; German.
                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 PLSSIFSRIGDP
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12; Conserv
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                               Similarity
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                                                                                                                215
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  100.0%; Score 61; DB 21; ilarity 100.0%; Pred. No. 0.00086; Conservative 0; Mismatches 0;
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Pred. No. 0.00068;
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                                                                  Length 215;
                 Indels
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                    Gaps
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AAR27471
ID AAR277
XX Hepath
KW Vacc:
XX Hepath
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XX W Vacc:
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XX Key
FT Regii

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The sequence given is encoded by an expression cassette which cc consists of the hepatitis B virus (HBV) M protein (small pre-S CC antigen, spaAg) gene precisely linked to a modified synthetic creation of the construction of a NYVAC recombinant expressing the HBV gene. CC the construction of a NYVAC recombinant expressing the HBV gene. CC ther HBV genes were also used in the construction. These were HBV CC ther HBV gene (S12/core, S1 + S2). Each of these gene sequences cc entire pre-S region (S12/core, S1 + S2). Each of these gene sequences were inserted individually into three different sites of NYVAC cc separated by from each other by large regions of vaccine strain of containing essential genes. NYVAC is a Copenhagen vaccine strain of containing virus which has been modified by deletion of six non-essential cc regions of the genome encoding known or potential virulence factors. The deletion loci were engineered as recipient loci for the insertion of foreign genes. The spacing of the three inserted sequences ensured cc that any recombination that did occur would lead to disruption of the vaccinia genome and would cause unviable vaccinia virus. See also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR27471 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR27471
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11-JUN-1991;
06-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis B virus; HBV; M protein; small pre-S antigen; spsAg; vaccinia virus; H6; early/late; promoter; NYVAC; recombinant; HBV L; large pre-S antigen; lpsAg; fusion protein; pre-S region; S12/core; S1; S2; Copenhagen vaccine strain; vaccinia virus; virulence factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     apaAg protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             deletion loci;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vaccine comprises recombinant, attenuated vaccinating against viral infections such HIV, HSV, EBV, CMV, mumps etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Riviere
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cox WI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig 9; 456pp; English.
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KJ, Norton EK,
M, Tartaglia J,
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92US-0847951.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9108-0666056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92WO-US01906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /label= S2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Francis J, Gettig RR, Jo
c, Paoletti E, Perkus ME,
J, Taylor J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     281
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pox-virus - use for as rabies, hepatitis B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Johnson
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This sequence is the hepatitis B virus (HBV) surface antigen (HBSAg)
CC pres2-S region. This sequence was used to create a recombinant antigen
CC library. The library comprises recombinant nucleic acids encoding
CC antigenic polypeptides and is produced by recombination of at least two
CC forms of nucleic acid, differing by at least two nucleotides, encoding
CC disease-related antigenic polypeptide. The library can be used to produce
CC a recombinant multivalent antigenic polypeptides of the invention, that
CC contains at least two antigenic polypeptides from different
CC contains at least two antigenic polypeptides are used in vaccines
CC infectious. The multivalent antigenic polypeptides are used in vaccines
CC infectious agents (bacteria, viruses, parasites, including Plasmodium
CC infectious agents (bacteria, viruses, parasites, including Plasmodium
CC arthritis, diabetes, multiple sclerosis); other inflammatory conditions
CC and cancer, also, where directed against sperm antigens, they can be used
CC optimised immune response against a wide variety of antigens, including attrains of a
CC particularly a broad spectrum response to many different strains of a
CC nathogen, including attrains that are likely to amount in that firely a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 13
AAY32835
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Best Local
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11-FEB-1998;
11-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                         Recombinant multivalent antigenic polypeptide produced recombining nucleic acid sequences and screening, used
                                                                                                                                                                                                                                                                                                                 Example 14; Fig 17; 153pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Вавв SH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MAXY-) MAXYGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HBsAg; PreS2-S; recombinant antigen library; disease-related antigen multivalent antigenic polypeptide production; infection; allergen; asthma; autoimmune disease; rheumatoid arthritis; diabetes; therapy; multiple sclerosis; inflammatory condition; cancer; contraception; immune response; hepatitis b surface antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9941383-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatitis b virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PreS2-S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12;
                                                                                                                                                                                                                                                                                                                                                   e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                AAZ10968.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Howard R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             281 AA;
                                                                                                                                                                                                                                                                                                                                                 infections and
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ilarity 100.0%;
Conservative (
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98US-0021769.
98US-0074294.
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Pred. No. 0.0011;
                     are likely to appear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stemmer WPC,
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                     in the future
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RESULT 14
AAR27472
ID AAR27
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Best Local S
Matches 12
  The sequence given is encoded by an expression vector which comprise the hepatitis B virus (HBV) L protein (large pre-S antigen, lpsAg) gene linked to the cowpox hemorraghic region (u) promoter. This DNN sequence was used in the construction of a NYVAC recombinant expressing the HBV gene. Other HBV genes were also used in the construction. These were HBV M protein (small pre-S antigen, spsAg) and a fusion protein composed of the entire pre-S region (S12/core, into three different sites of NYVAC separated by from each other by large regions of vaccinia DNA containing essential genes. NYVAC is copenhagen vaccine strain of vaccinia virus which has been modified deletion of six non-essential regions of the genome encoding known o
                                                                                                                                                                                                                                                                                                                                                            07-MAR-1991;
11-JUN-1991;
06-MAR-1992;
                                                                                                                                                              Disclosure; Fig 11; 456pp; English
                                                                                                                                                                                     Vaccine comprises recombinant, attenuated pox-virus - use for vaccinating against viral infections such as rabies, hepatitis B, HIV, HSV, EBV, CMV, mumps etc.
                                                                                                                                                                                                                                              N-PSDB; AAQ29104.
                                                                                                                                                                                                                                                                                                            Cox WI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis B virus; HBV; M protein; small pre-S antigen; spsAg; vaccinia virus; hemorraghic region; u; promoter; NYVAC; recombinant; HBV L; large pre-S antigen; lpsAg; fusion protein; pre-S region; $12/core; $1; $2; Copenhagen vaccine strain; vaccinia virus; virulence factor; deletion loci; recipient loci.
                                                                                                                                                                                                                                                                                                Limbach
                                                                                                                                                                                                                                                                                                                                                                                                              09-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
region
                                                                                                                                                                                                                                                                                                                                   (VIRO-) VIROGENETICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    misc_RNA
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                                                                                                                                                                                                                                                        1992-331718/40.
                                                                                                                                                                                                                                                                                KJ, Norton El
M, Tartaglia
                                                                                                                                                                                                                                                                                              De Taisne C, |
KJ, Norton EK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; Protein; 389 AA.
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91US-0713967.
92US-0847951.
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/label= S2
164..389
/label= S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cocation/Qualifiers
                                                                                                                                                                                                                                                                           Francis J, Get
K, Paoletti E,
J, Taylor J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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Pred. No. 0.0011;
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                                                                                                                                                                                                                                                                                          Gettig RR, J
8, Perkus MB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20;
                                                                                                                                                                                                                                                                                                      Johnson GP
                                                                                                                                                                                                                                                                                          Pincus SE;
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                                                                                                           lpsAg)
This DNA
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regions of the genome

encoding

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RESULT 15
AAR27474
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             potential virulence factors. The deletion loci were engineered as recipient loci for the insertion of foreign genes. The spacing of the three inserted sequences ensured that any recombination that did occur would lead to disruption of the vaccinia genome and would cause unviable vaccinia virus. See also AAQ35501-864.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                           07-MAR-1991;
11-JUN-1991;
06-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis B virus; HBV; M protein; small pre-S antigen; spsAg; vaccinia virus; Amsacta moorei entomopoxvirus; AmsPV; 42 kD; pro NYVAC; recombinant; HBV L; large pre-S antigen; lpsAg; fusion pre-S region; S12/Coore; S1; S2; Copenhagen vaccine strain; vaccinia virus; virulence factor; deletion loci; recipient loci.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                    Region
                                                                                                                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    149 PLSSIFSRIGDP 160
The sequence given is encoded by an expression cassette which comprises the haparitis B virus (HBV) L protein (large pre-S antigen, lpsAg) gene which is precisely linked to the Amsacta moorei entomopoxytrus (AmEPV) 42 kD promoter. This DNA sequence was used in the construction of a NYVAC recombinant expressing the HBV gene.
                                                                                        Vaccine comprises recombinant, attenuated vaccinating against viral infections such HIV, HSV, EBV, CMV, mumps etc.
                                                                                                                                                                                               COX WI,
                                                                                                                                                                                                                                                                                        09-MAR-1992;
                                                                                                                                                                                                                                                                                                              17-SEP-1992
                                                                                                                                                                                                                                                                                                                                    WO9215672-A
                                                                                                                                                                                                                                                                                                                                                                                                                 Region
                                                                                                                                       N-PSDB; AAQ29106
                                                                                                                                                                                                                      (VIRO-) VIROGENETICS
                                                                    Disclosure; Fig 15; 456pp; English.
                                                                                                                                                                         Riviere
                                                                                                                                                                                    Limbach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PLSSIFSRIGDP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         promoter/S12/core gene.
                                                                                                                                                     1992-331718/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                      De Taisne C, Fru-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            389 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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91US-0713967.
92US-0847951.
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109..163
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164..389
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                                                                                                                                                                                                                       CORP
                                                                                                                                                                         Francis J,
K, Paoletti
J, Taylor J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               389 AA.
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Pred. No. 0.0016;
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                                                                                                                                                                       'n,
                                                                                                                                                                                       Gettig RR, Jo
E, Perkus ME,
                                                                                                          pox-virus - use for
as rabies, hepatitis
                                                                                                                                                                                        Johnson G
E, Pincus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 389;
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                                                                                                                                   Query Match
Best Local S
Matches 12
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                                                                                                                                                                                                                                                       vaccinia genome and would AAQ35501-864.
                                                                                                                                                                                                                   Sequence
                                                     149
                                                                                   1 PLSSIFSRIGDP 12
                                                     PLSSIFSRIGDP 160
                                                                                                                          n 100.0%;
Similarity 100.0%;
12; Conservative 0.
                                                                                                                                                                                                                       389
2003, 09:23:52
                                                                                                                                         0
                                                                                                                                                             Score 61; DB 13;
Pred. No. 0.0016;
                                                                                                                                              Mismatches
                                                                                                                                                                                  DB 13;
                                                                                                                                                                                  Length 389;
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                                                                                                                                                Gaps
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Search completed: February Job time : 2.92335 secs

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Result
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Post-processing: Minimum Match 0% Maximum Match 100%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Title:
Perfect score:
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                                                                                                                                       Score
             protein search, using sw model
                                                                                                                                     Query
Match
                         100.0
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902.626 Million cell updates/sec
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|: /cgn2_6/ptodata/1/paa/
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/ Cgm2_6/ptodata/1/paa/USO6_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO6_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO6_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO61_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO62_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO63_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO63_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO63_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO66_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO66_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO66_COMB.pep: *
/ Cgm2_6/ptodata/1/paa/USO69_COMB.pep: *
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US-09-830-981-16

US-09-830-981-16
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	Sequence 106, App	100	106	Seguence 106	e 106	equence 106	D (106	106	•	106.	106	106	106	106	106	116	217.	equence 214.	equence 219	quence 216,	equence 12.		equence 12	e 12 App	-	213	בקקא יי פ	0		sequence 45,	; -	segmence au, appli	- t	2 1	3.	و م	2		

STATEMENT'S

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US-09-830-981-2

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Sequence 2, Application US/09830981
GENERAL INFORMATION:
APPLICANT: Eberhard Hildt
APPLICANT: Etephanie Schmidt
TITLE OF INVENTION: Polypeptide Mediating Ce.
FILE REFERENCE: 033392-001
CURRENT APPLICATION UNMER: US/09/830,981
CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION NUMBER: PCT/DE99/03506
PRIOR FILING DATE: 1999-11-03
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR PILING DATE: 1999-11-03
PRIOR PILING DATE: 1999-11-03
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
TENGTU. 12
                                                                                                        US-09-830-981-2
       Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative 0
                                                                                                               LENGTH: 12

TYPB: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description
OTHER INFORMATION: Cell permeab
                                                                                                                 Description of Artificial Sequence:
Cell permeability mediating polypeptide
            0
    Score 61; DB 22;
Pred. No. 0.00042;
), Mismatches 0;
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                                                Length 12;
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Gaps
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PLSSIFSRIGDP 12

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RESULT 2

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PRIOR APPLICATION NUMBER: POT/DE99/03506
PRIOR FILING DATE: 1999-11-03
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR FILING DATE: 1998-11-03
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
LENGTH: 12
TYPE: PRT
ORGANISM: Hepadnavirus
US-09-830-981-4
                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/830,981
CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION NUMBER: PCT/DE99/03506
PRIOR FILING DATE: 1999-11-03
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR FILING DATE: 1998-11-03
NUMBER OF SEQ ID NOS: 20
SOPTWARE: PATENTIN Ver. 2.1
SEQ ID NO 12
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                                                                     RESULT 4
US-09-830-981-14
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US-09-830-981-12
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GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: Stephanie Schmidt
TITLE OF INVENTION: Polypeptide Mediating Cell Permeability
FILE REFERENCE: 033392-001
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                                Sequence 14, Application US/09830981
GENERAL INFORMATION:
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APPLICANT: Stephanie Schmidt
TITLE OF INVENTION: Polypeptide Mediating Cell Permeability
FILE REPERENCE: 033392-001
 APPLICANT:
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Local Similarity 100.0%;
es 12; Conservative 0
                                                                                                                                              atch 100.0%; cal Similarity 100.0%; 12; Conservative 0;
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Eberhard Hildt
Stephanie Schmidt
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Pred. No. 0.00042;
Mismatches 0;
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Sequence 16, Application US/09830981
GENERAL INFORMATION:
APPLICANT: Eberhard Hildt
APPLICANT: Stephanie Schmidt
TITLE OF INVENTION: Polypeptide Mediating Ce.
FILE REFERENCE: 033392-001
CURRENT APPLICATION NUMBER: US/09/830,981
CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION NUMBER: PCT/DE99/03506
PRIOR PILING DATE: 1999-11-03
PRIOR FILING DATE: 1999-11-03
Sequence 18, Application US/09830981
GENERAL INFORMATION:
APPLICANT: Eberhard Hildt
APPLICANT: Eberhard schmidt
TITLE OF INVENTION: Polypeptide Mediating Ce.
FILE REFERENCE: 033392-001
CURRENT APPLICATION UNWBER: US/09/830,981
CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION NUMBER: PCT/DE99/03506
PRIOR FILING DATE: 1999-11-03
PRIOR FILING DATE: 1999-11-03
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 18
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US-09-830-981-14
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CURRENT APPLICATION NUMBER: US/09/830,981
CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION NUMBER: PCT/DE99/03506
PRIOR FILING DATE: 1999-11-03
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR FILING DATE: 1998-11-03
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 14
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US-09-830-981-16
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US-09-830-981-18
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Best Local Similarity
Matches 12; Conserv
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NUMBER OF SEQ ID NOS: 20
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 16
LENGTH: 12
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Best Local Similarity 100.0%;
Matches 12; Conservative (
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100.0%; Pred. No. 0.00042;
tive 0; Mismatches 0;
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Pred. No. 0.00042;
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APPLICANT: Hidt, Eberhard
APPLICANT: Hofschneider, Peter
ITILE OF INVENTION: Particles for Gene Therapy
FILE REFERENCE: 107070-120 (VOS-013)
CURRENT APPLICATION NUMBER: US/09/890,752A
CURRENT FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: PCT/DE00/00363
PRIOR APPLICATION NUMBER: DE 199 04-800.2
PRIOR FILING DATE: 2000-02-04
PRIOR FILING DATE: 1999-02-05
NUMBER OF SEQ ID NOS: 21
SOPTWARE: Patentin version 3.1
SEQ ID NO 20
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
FEATURE:
FEATURE:
TYPE: THEORMATTON: cell permeability-mediating
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                                                                                      US-09-890-752A-20
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US-09-830-981-20
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US-09-830-981-18
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                          Query Match
Best Local Similarity
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SEQ ID NO 20
LENGTH: 12
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CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION NUMBER: PCT/DE99/03506
PRIOR FILING DATE: 1999-11-03
PRIOR APPLICATION NUMBER: DE 198 50 718.6
PRIOR FILING DATE: 1998-11-03
NUMBER OF SEQ ID NOS: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Eberhard Hildt
APPLICANT: Stephanie Schmidt
TITLE OF INVENTION: Polypeptide Mediating Cell Permeability
FILE REFERENCE: 033392-001
                                                                                          OTHER INFORMATION: cell permeability-mediating peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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12; Conserv
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  100.0%; Score 61; DB 22;
100.0%; Pred. No. 0.00042;
tive 0; Mismatches 0;
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US-10-226-956-284
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US-10-077-555-9
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                                                                                 Sequence 284, Application US/10226956 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 12; Conserv
            APPLICANT: Brophy, Colleen
APPLICANT: Komalavilas, Pac
APPLICANT: Panitch, Alyssa
APPLICANT: Joshi, Lokesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Hwu , Paul L.
TITLE OF INVENTION: A NEW FUSION PROTEIN FOR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: GeneShuttle Biopharm, Inc. APPLICANT: Hwu , Paul L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Wang, Rong-fu
TITLE OF INVENTION: Use of Cell-Penetrating Peptides to Generate Antitumor Immunity
FILE REFERENCE: P02373US1/10200806
CURRENT APPLICATION NUMBER: 102/0/77,555
CURRENT FILING DATE: 2002-02-15
CURRENT FILING DATE: 2002-02-15
                                                                                                                                                                                                                                                                                                                    FEATURE:
NAME/KEY: MISC FEATURE
OTHER INFORMATION: The HBV PreS2 antigen consisting of the translocation motif from OTHER INFORMATION: residues 41-52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/144,549
CURRENT FILING DATE: 2002-05-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: MBHB 02-340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/268,687 PRIOR FILING DATE: 2001-02-15
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                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Hepatitis B virus
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ORGANISM: Hepatitis B virus
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                                                                                                                                                                                                                                                                  100.0%; Score 61; DB 25; 100.0%; Pred. No. 0.00042;
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100.0%; Pred. No.
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PRIOR FILING DATE: 2001-08-23
NUMBER OF SEQ ID NOS: 320
SOFTWARE: PALENTIN VERSION 3.1
SEQ ID NO 284
LENGTH: 12
US-09-021-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Smith, Timothy L.
REGISTRATION NUMBER: 35,367
REFERENCE/DOCKET NUMBER: 0180
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 45:
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/226,956
CURRENT FILING DATE: 2002-08-23
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PALENTION DATA:

APPLICATION NUMBER: US/09/021,769

FILING DATE: 11-FEB-1998

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APPLICANT: Howard, Russell
TITLE OF INVENTION: Methods and Compositions for Vaccine
TITLE OF INVENTION: Engineering
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 102 amino acid
                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: peptide
                   NAME/KEY: Peptide LOCATION: 1..102 OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION:
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|| Similarity 100.0%;
| 12; Conservative 0
69-45
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Stemmer, Willem P.C.
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                       /note= "truncated Hepatitis B surface
antigen (HBsAg) (PreS2 plus S regions)
polypeptide"
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pred. No. 0.00042;
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US-09-248-716-6
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US-09-248-716-6
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Best Local Similarity
Matches 12; Conserve
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US-09-724-890-45
; Sequence 45, Application US/09724890
; GENERAL INFORMATION:
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GENERAL INFORMATION:
APPLICANT: Punnone
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Best Local Similarity
Matches 12; Conserv
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LENGTH: 102
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APPLICANT: Howard, Russell
APPLICANT: Maxygen, Inc.
APPLICANT: Maxygen, Inc.
TITLE OF INVENTION: Optimization of Immunomodulatory Properties of Genetic TITLE OF INVENTION: Vaccines
FILE REFERENCE: 018097-030300US
CURRENT APPLICATION NUMBER: US/09/248,716
CURRENT FILING DATE: 1999-02-10
PRIOR APPLICATION NUMBER: US 60/074,294
PRIOR FILING DATE: 1998-02-11
NUMBER OF SEG ID NOS: 02-11
NUMBER OF SEG ID NOS: 03-0-11
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION MOMBER: US/09/724,890
FILING DATE: 28-Nov-2000
                                                                                                                                                                                                                                                                                                                                              APPLICANT: Punnonen, Juha
Stemmer, Willem P.C.
Whalen, Robert Gerald
Howard, Russell
                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
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                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Methods and Compositions for Vaccine
Engineering
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                                                                                                                                                                                               CITY: San Francisco
STATE: California
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                                                                                                                                                         ZIP: 94111-3834
                                                                                                                                                                             COUNTRY: USA
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Pred. No. 0.0044;
); Mismatches 0
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Pred. No.
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                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence:Plant-optimized; OTHER INFORMATION: pre S1/S2.
PCT-US99-31020-41
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PCT-US99-31020-41
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Matches
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GENERAL INFORMATION:
APPLICANT: Boyce Thompson Inst. for Plant Res. at Cornell
TITLE OF INVENTION: Expression of Immunogenic Hepatitis B Surface Antigens
TITLE OF INVENTION: in Transgenic Plants
FILE REFERENCE: 4868/85428
CURRENT APPLICATION NUMBER: PCT/US99/31020
CURRENT APPLICATION NUMBER: PCT/US99/31020
CURRENT FILLING DATE: 2000-04-10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 41
LENGTH: 164
TYPE: PRT
                                                                                                                                                                                                            Query Match
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41 PLSSIFSRIGDP 52
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PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/021,769

FILING DATE: 11-FEB-1998

ATTORNEY/AGENT INFORMATION:

NAME: Smith, Timothy L.

REGISTRATION NUMBER: 35,367

REFERENCE/DOCKET NUMBER: 018097-021500US

TELEPHONE: (415) 576-0200

TELEPAY. (415) 576-0200
                                                                                                               1 PLSSIFSRIGDP 12
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TYPE: amino acid
STRANDEDNESS: <Unknown>
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ALIGNMENTS

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AC 091IC1;
DT 01-DBC-2001 (TrEMBLrel. 19, Created)
DT 01-DBC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-DBC-2001 (TrEMBLrel. 20, Last annotation update)
DT 01-MAR-2002 (TrEMBLRel. 20, Last sequence update)
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Middle surface antigen.
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EMBL; AF136461; AAD44711.1; -.
InterPro: IPR000349; Hepvir_surfAg.
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EMBL; U87737; AAC58018.1; -.
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EMBL; AJ34417; CAC87014.1; -.

ENGL; AJ34417; CAC87014.1; -.
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Submitted (MAR-1995) to the EMBL/GenBank/DDBU databases.
EMBL; X85254; CAA59510.1;
InterPro; IPR000349; Hepvir surfAg.
Pfam; PF00695; VMSA; 1.
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Viruses; Retroid viruses; Hepadnaviridae;
NCBI_TaxID=10407;
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Q67954;
Q1-NOV-1996 (TrEMBLrel. 01,
Q1-NOV-1996 (TrEMBLrel. 01,
Q1-MAR-2002 (TrEMBLrel. 20,
                                                                                                                Laī M.B., Mazzoleni A.P., Balestrieri A.;
"Sequence analysis of HBV genomes isolated fro
chronic liver disease.";
Submitted (JAN-1994) to the EMBL/GenBank/DDBJ
EMBL, X77309; CAA54515.1;
                                                                                                                                                                                                                                                                                                                                  HBV surface proteins (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae;
NCBI TaxID=10407;
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                                                             InterPro; IPR000349; Hepvir_surfAg
Pfam; PF00695; vMSA; 1.
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"Sequence analysis of pre-S/S gene of hepatitis B genetypes A. D. and F isolated in Brazil.";
Arch. Virol. 141:1767-1773(1996).
EMBL, U55227, AAB4/473.1;
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MEDLINE=97049068; PubMed=8893798;
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Moraes M.T., Gomes S.A., Niel C.;
"Sequence analysis of pre-S/S gene of hepatitis genotypes R. D. and F isolated in Brazil.";
Arch. Virol. 141:1767-1773(1996).
EMBL; U55224; AAB47470.1;
InterPro; IPR000349; Hepvir_surfag.
Pfam; PF006595; VMSA; 1.
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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Pred. No. 0.0015;
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Pfam; PF00695;
CHAIN 164
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01-NOV-1999 (TrEMBLrel.
01-NOV-1999 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
                                          SEQUENCE
                                                                     Submitted (JAN-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ131956; CAA10540.1; ...
InterPro; IPR000349; Hepvir surfAg.
Pfam; PF00695; vMSA; 1
                                                                                                                                        Petzold D.R.;
Submitted (JA
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01-NOV-1998
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01-MAR-2002
                                                                                                                                                                               SEQUENCE FROM N.A.
STRAIN=1;
                                                                                                                                                                                                                          Petzold D.R., Tautz B., Wolf F., Drescher J.; "Infection chains and evolution rates of Hepatitis B virus in cardiac transplant recipients infected nosocommially."; J. Med. Virol. 58:1-10(1999).
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STRAIN=1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis B virus.
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EMBL; AJ003116; CAA05872.1; -.
                                                                                                                                                                                                                                                                                                                MEDLINE=99238050; PubMed=10223539;
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Viruses, Retroid viruses,
NCBI TaxID=10407;
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Q91C52;
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01-DEC-2001
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01-MAR-2002
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01-DEC-2001 (TYEMBLIE). 19, Last sequence update)
01-DEC-2002 (TYEMBLIE). 20, Last annotation update)
101-WAR-2002 (TYEMBLIE). 20, Last annotation update)
Large/middle/small S proteins.
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submitted (AUG-2000) to the EMBL/GenBank/DDBJ
EMBL; AF297619; AAK97179.1; -.
InterPro; IPR000349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
SEQUENCE 389 AA; 42580 MW; F4336E04992CE6
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J. Med. Virol. 64:441-454 (2001).
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus
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Hepatlis B virus.

Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.

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                                                                                                                                                                                                                                        MEDLINE=21361467; PubMed=1146872B; Owiredu W.K., Kramvia A., Kew M.C.; Owiredu W.K., Kramvia A., Kew M.C.; Phepatitis B virus Dan in serum of healthy black African adults positive for hepatitis B surface antibody alone; possible association with recombination between genotypes A and D."; J. Med. Virol. 64:441-454 (2001).
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Owiredu W.K.B.A., Kramvis A.,
Submitted (AUG-2000) to the E
EMBL, AF297620; AAK97183.1;
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(TrEMBLrel. 19, Last sequence update)
(TrEMBLrel. 20, Last annotation update)
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                                                                   Kew M.C.;
EMBL/GenBank/DDBJ
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Pfam; PP00695; vMSA; 1.
SEQUENCE 389 AA; 42937 MW; 1709
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"Sequence analysis of HBV genomes isolated
negative chronic liver disease.";
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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01-JUN-2002 (TrEMBLrel. 21,
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Viruses; Retroid v
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NCBI_TaxID=10407;
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                                                                                           Hepatitis B surface antigen-negative Hepatology 0:0-0(0).
EMBL; AJ344117; CAC87013.1;
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RESULT 15
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SEQUENCE FROM N.A.

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STRALN=702-4;

Huangfu J., Dong J., Deng H.;

"Preliminary Study on PreS2 Region of Hepatitis B Virus.";

Submitted (JUN-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AF393221; AAX84360.1; -.

InterPro; IPR000349; Hepvir_surfAg.

Pfam; PF00695; vMSA; 1.

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49
SEQUENCE 49 AA; 5045 MW; 5CA3FD6A356B59C3 CRC64;
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O91IC3;
O1-DEC-2001 (TrEMBLrel. 19, Created)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Pre-S2 protein (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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7: sp_mhc:*
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		۴			SUMMARIES	
Result No.	Score	Query Match	Length	BG	ID	Description
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10	1875	97.2	389	12	Q8QXQ3	hepatiti
11	1875		389	12	Q8QXP9	henderiti
12	1874	97.1	389	12	Q9QAF2	Oggaf2 henatitis
13	1864	96.6	686	12	Q9QAF9	9 henatiti
14	1864	96.6	389	12	O9WNS1	- '
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Q96843,
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Submitted (AUG-1996) to the EMBL/GenBank/DDBJ
EMBL; Y07587; CAA68861.1; -.
InterPro; IPR000349; Hepvir_surfAg.
Pfam; PP00695; VMSA; 1.
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Stoll-Becker S., Repp R., Glebe D., Schaefer
Lampert F., Gerlich W.H.;
"Transcription of hepatitis B virus in peripl
cells from persistently infected patients.";
J. Virol. 71:5399-5407(1997).
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Viruses; Retroid viruses; H
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Hannoun C., Horal P., Lindh M.;
Hubmitted (JAN-1999) to the EMBL/Geni
EMBL; AF121241; AAF24674.1; -.
EMBL; AF121240; AAF24667.1; -.
EMBL; AF121240; AAF24667.1; -.
InterPro; IPR000349; Hepvir_BurfAg.
Pfam; PF00695; vMSA; 1.
SEQUENCE 389 AA; 42814 MW; CD1891
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Q1-MAY-2000 (TrEMBLrel. 1
Q1-MAY-2000 (TrEMBLrel. 1
Q1-MAR-2002 (TrEMBLrel. 2
                                                                                  Q67871
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Q1-NOV-1996 (TrEMBLrel. Q1,
Q1-NOV-1996 (TrEMBLrel. Q1,
Q1-MAR-2002 (TrEMBLrel. 20,
Pre S1/pre S2/S ORF.
Hepatitis B virus.
Viruses; Retroid viruses; H
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STRAIN-HBV/95-1758, AND HBV/94-11066;
MEDLINE=20109034; PubMed=10640544;
Hannoun C., Horal P., Lindh M.;
"Long-term mutation rates in the hepatitis
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus
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Pred. No. 4.2e-153;
1; Mismatches 2;
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Best Local Similarity
Matches 337; Conserv
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01-NOV-1996 (TrEMBLrel. 0:
01-NOV-1996 (TrEMBLrel. 0:
01-MAR-2002 (TrEMBLrel. 2:
                                                                                                                                                                                            SEQUENCE FROM N.A.

Karaylannis P.;

Submitted (MAY-1996) to the EMBL/GenBank/DDBJ

EMBL; X97848; CAA66425.1; -.

InterPro; IPRO0349; Hepvir_surfag.

Pfam; PF00695; vMSA; 1.

SEQUENCE 389 AA; 42717 MW; 495EF28627FBF62
                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDINE-97025699; PubMed=8871878;
Alexopoulou A., Karayiannis P., Hadziyannis S.J.;
"Whole genome analysis of hepatitis B virus from 4 cases with fulminant hepatitis.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis B virus.
Viruses, Retroid viruses, Hepadnaviridae,
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EMBL; X65257; CAA46349.1; "InterPro; IPR000349; Hepvir_surfAg.

Pfam; PP00695; vMSA; 1.
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                                                                                               Score 1883; DB 12;
Pred. No. 6.2e-153;
3; Mismatches 2;
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Pred. No. 6.2e
2; Mismatches
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Q67920;
Q1-NOV-1996
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Q1-MAR-2002
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Hepatitis B virus.
Viruses, Retroid viruses,
NCBI TaxID=10407;
                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=97025699; PubMed=8871878;
Alexopoulou A., Karaylannis P., Hadziyannis
"Whole genome analysis of hepatitis B virus
fulminant hepatitis.";
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STRAIN=PRE-CORE VARIANT;
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TTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAPGKFLWEWASARFSWI
            TTPAQGTSMYESCCCTXPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                    CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLDVCFLIPGSSTTSTGPCRTC
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                                                                                                    TSGFLGPLLVLQAGFFLLTRILTIPOSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
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Last annotation updat
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Pred. No. 1.4e-152;
2; Mismatches 3;
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Best Local S
Matches 337
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STRAIN=GENOTYPE D;

MEDLINE=89010694; PubMed=3171552;

MEDLINE=89010694; Nashtypes H, Sastrosoewignjo R.I., I

MIYAKAWA Y., MAYUMI M.;

MIYAKAWA Y., MAYUMI M.;

MIYAKAWA Y., MAYUMI M.;

STOMPATION OF SURVEY DE STRAIN OF SEQUENCE FROM N.A.;

J. Gen. Virol. 69:2575-2583(1988).

J. Gen. Virol. 69:2575-2583(1988).

L. J. Gen. Virol. 69:2575-2583(1988).

R. EMBL; AB033559; BAA85378.1; -.

R. EMBL; AB033559; BAA85378.1; -.

R. EMBL; AB033559; BAA85378.1; -.

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Q9QMH6;
01-MAY-2000
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PRES1/PRES2/S.

Hepatitis B virus.

Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.

NCBI TaxID=10407;
Q91F39;
Q91F39;
01-OCT-2000
01-OCT-2000
01-MAR-2002
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Matches 336; Conserv
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"First Chinese HBV strain: genotype D complete sequence report.";
Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.
EMBL, AF280817; AAF81508.1; -.
InterPro; IPR000349; Hepvir_surfAg.
Pf4mm; PF00695; VMSA; 1.
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Viruses; Retroid viruses;
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Viruses; Retroid viruses;
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                                                                                                                  "Molecular cloning and sequencing of two complisolates of Human Hepatitis B Virus.";
Submitted (AUG-1994) to the EMBL/GenBank/DDBJ EMBL; 235716; CARB4788:1; -.
Interpro; IPR000349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
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CHAIN
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Pred. No. 2.
      <u>ب</u>
      Score 1875; DB 12;
Pred. No. 3e-152;
2; Mismatches 4;
                                                                                    SURFACE PROTEIN S. 5D27F0BA97A825D5
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Q8QXQ3;
01-JUN-2002
01-JUN-2002
01-JUN-2002
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Jeantet D., Chemin I., Mandrand B., Zoulim F., Trepo C., Kay A.;
"Characterization of two Hepatitis B virus populations isolated;
Hepatitis B surface antigen-negative patient.";
                                                                                                                                                                                                                                              Hepatology 0:0-0(0).
EMBL; AJ344116; CAC87019.1; -.
SEQUENCE 389 AA; 42846 MW;
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STRAIN=HBSAG NEGATIVE
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Viruses; Retroid viruses;
NCB TaxID=10407;
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98.5%;
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Q9QAF2;
01-MAY-2000 (TrEMBLrel. 13, 0
01-MAY-2000 (TrEMBLrel. 13, I
01-MAR-2002 (TrEMBLrel. 20, I
Large S protein.
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Viruses; R
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Q8QXP9;
01-JUN-2002 (TIEMBLrel. 21, C
01-JUN-2002 (TIEMBLrel. 21, L
01-JUN-2002 (TIEMBLrel. 21, L
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EMBL; AJ344117; CAC87013.1; -.
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Viruses, Retroid viruses,
NCBI TaxID=10407,
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Retroid v
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Hepadnaviridae; Orthohepadnavirus.
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Mismatches 3;
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DRIE REPRESENTATION OF THE PROPERTY OF THE PRO
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Hannoun C., Horal P., Lindh M.;
Hannoun C. (JAN-1999) to the EMBL/GenBank/DDBJ
Submitted (JAN-1999) to the EMBL/FenBank/DDBJ
EMBL; AF121242; AAF24681.1;
InterPro; IPR000349; Hepvir_BurfAg.
Pfam; PF00695; VMSA; 1.
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                                                                                                                                                                                                               STRAIN-HBV/1303;
WEDLINE=20109034; PubMed=10640544;
Hannoun C., Horal P., Lindh M.;
"Long-term mutation rates in the horal process of t
Hannoun C., Horal P., Lindh M.;
Submitted (JAN-1999) to the EMBL/GenBank/DDBJ
EMBL; AF121239; AAF24661.1;
InterPro; IFR000349; Hepvir_surfAg.
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01-MAR-2002 (TrEMBLrel.
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Viruses; Retroid viruses;
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SEQUENCE 389 AA; 42890 MW;
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01-NOV-1999 (TrEMBLrel. 12,
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Gerner P., Lausch
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                                                                                                                                                                       Signal.
                                                                                                                                                                                                                                                 Submitted (MAY-1999) to the EMBL/GenBank/DDBJ EMBL; AF151735; AAD41361.1; -
                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                 "Hepatitis B virus core promoter mutations in children with multiple anti-HBe/HBeAg reactivations result in enhanced promoter activity."; J. Med. Virol. 59:415-423(1999).
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B., Friedt M., Tratzmuller R.,
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Pred. No. 2.
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S PROTEIN.
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                                                                                                                                                                                                                                                                                                                                                                    Ouery Match 96.5
Hest Local Similarity 97.4
Matches 333; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gunther S., Wilma P., Meisel H., Will H.;
Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF043594; AAC40811.1; -.
InterPro; IPR000349; Hepvir_BurfAg.
Pfam; PF00695; vMSA; 1.
CHAIN
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389
SPOTEIN.
SBQUENCE 389 AA; 42795 MW; 3369787D78F188F9 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDINE-98240943; PubMed=9581787;
Gunther S., Paulij W., Meisel H., Will H.;
"Analysis of hepatitis B virus populations in
treated patient reveals predominant mutations
Changing e-antigenicity.";
Virology 244:146-160(1998).
186 CPPTCPGYRWMCLRREIIFLEILLLCLIFLLVLLDYQGMLFVCFLIPGSSTTSTGFCRTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=1/91;
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01-NOV-1998 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
                                                                                           168
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI TaxiD=10407;
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01-NOV-1998
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                                                                      TSGLLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
                                                                                                                                                                 AMOWNSTTFHOTLODPRVRGLYFPAGGSSSGTVNPVPTTASPISSIFSRIGDPALNNENI 167
                                                                                                                                                                                                  AMQMUSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSDISSIFSRIGDPALNMENI 125
                                                                                             TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
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                                                                                                                                                                                                                                                                                                                                                             96.5%; Score 1861; DB 12;
97.4%; Pred. No. 4.7e-151;
tive 3; Mismatches 6;
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the C-gene and
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Search completed: February 3, 2003, 09:25:43
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Listing first 45 summaries
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              Score
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Match Length
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"Nucleotide sequence of the hepatitis
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"Subtype aww variant of hepatitis i
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Pred. No. 2.3e-125;
2; Mismatches 4;
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Best Local Similarity
Matches 332; Conserv
                                                                                                       This SWISS-PROT entry is copyright. It is produced through a cobetween the Swiss Institute of Bioinformatics and the EMBL of the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and for entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
      CHAIN
CARBOHYD
                                                                            EMBL; V00867; CAA24234.1; ALT_INIT.
PIR; A03705; SAVLA.
                                                                                                                                                                                                                  MEDLINE=85107103; PubMed=3968537;
Takeshima H., Inokoshi J., Namiki M., Shimada J., Omura S.;
"Structural analysis of the gene coding for hepatitis B virus
antigen and its product.";
J. Gen. Virol. 66:195-200(1985).
                                                                                                                                                                                                                                                                                                     MEDIINE-83168919; PubMed-6300776;
Ono Y., Onda H., Sasada R., Igarashi K., S
"The complete nucleotide sequences of the
DNA; subtype adr and adw.";
Nucleic Acids Res. 11:1747-1757(1983).
                                                            InterPro; IPR000349; Hepvir surfAg
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                                PROPEP
                                            Antigen.
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Viruses; Retroid viruses; Hepadnaviridae;
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21-JUL-1986 (Rel. 01, Last seq
15-JUN-2002 (Rel. 41, Last ann
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Last annotation updat
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Pred. No. 3.7e-124;
2; Mismatches 8;
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Best Local Similarity
Matches 315; Conserv
          Antigen.
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                                                                                                                                                                   EMBL; D00220; BAA00159.1;
PIR; C28885; SAVLCP.
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01-OCT-1989 (Rel. 12, Created)
01-OCT-1989 (Rel. 12, Last sequence up
01-APR-1990 (Rel. 14, Last annotation
                                                                                                                             Pfam; PF00695; vMSA;
                                                                                                                                                                                                                                                                                                                                                                               "The complete nucleotide sequence of the genome of a hepatitis virus isolated from a naturally infected chimpanzee.";
J. Gen. Virol. 69:1383-1389(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Harrison
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NCBI_TaxID=10414;
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359 GLSPTVWLSVIWMMWYWGPSLYSILSPPLPLLPIFFCLWVYI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 surface antigen
                                                                                                                                                IPR000349;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B virus (strain Retroid viruses;
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164
4
112
166
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92.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                A.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lsh / chimpanzee isolate).
Hepadnaviridae; Orthohepadnavirus.
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    MAJOR SURFACE ANTIGEN.
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
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N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
, 16DB0B15C6CB697A CRC
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tsiquaye K.N., Zuckerman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
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on update)
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IN O1-UIL
DT 01-UIL
DT 15-UIL
DT 15-UIL
DT 01-UIL
DT 01-U
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Best Local Sim
Matches 313;
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q02317;
01-JUL-1993
01-JUL-1993
01-JUL-1993
                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                  isolated from a Philippino: Comparison with of the same subtype.";
J. Gastroenterol. Hepatol. 3:215-222(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis B virus (subtype adw / strain Philippino/pFDW294).
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=31514;
    CARBOHYD
CARBOHYD
CARBOHYD
CARBOHYD
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
Estacio R.C., Chavez C.C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Estacio R.C., Chavez C.C., Okamoto F Domingo E., Mayumi M.; "Nucleotide sequence of a hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        168
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                                                                                                                                PROPEP
                                                                                                                                                                             Pfam; .PF00695;
                                                                                                                                                                                                                        EMBL; M57663; AAA69680.1; -
                                                                                                                                                           Antigen.
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TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGAPVCLGQNSQSPTSNHSPTS
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                                                                                                                                                                             IPRO00349; Hepvir_surfAg
)695; vMSA; l.
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(Rel. 26, Last sequence update)
(Rel. 26, Last annotation updat
ce antigen precursor.
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91.5%;
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; 87D1F93382A10DBD CRC64;
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Pred. No. 2.3e-117;
                        MAJOR SURFACE ANTIGEN.
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
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Best Local S
Matches 311
Query Match
Best Local S
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01-OCT-1989 (Rel. 12, Created) 
01-OCT-1989 (Rel. 12, Last sequence up 
15-JUL-1998 (Rel. 36, Last annotation 
Major surface antigen precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                     HPBV4
                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its way by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                        VMSA_HPBV4
                               Antigen.
PROPEP
CHAIN
CARBOHYD
CARBOHYD
CARBOHYD
CARBOHYD
SEQUENCE
                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=10409;
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Viruses; Retroid viruses; Hepadn
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                                                                                                                                                                                                                                                                                        Fujiyama A.,
                                                                                                                                                                                                                                                                                         MEDLINE-83246570; PubMed=6306594;
Fujiyama A., Miyanohara A., Nozaki C.,
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EMBL; X01587; CAA25743.1; ALT_INIT.
                                                                                                                                                                                                                                                Nucleic Acids Res. 11:4601-4610(1983).
                                                                                                                                                                                                                                                                               Matsubara K.;
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                                                                                                                                                                       Antigen.
PROPEP
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PIR; G28925; SAVLJ1.
                                                                                                                                                                                                                                                                                                                                                                                                         MEDIJINE=89010694; PubMed=3171552; Chamboto H., Tsuda F., Sakugawa H., Sastrosoewignjo R Miyakawa Y., Mayumi M.; "Typing hepatitis B virus by homology in nucleotide comparison of surface antigen subtypes."; J. Gen. Virol. 69:2575-2583(1988).
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P17398;
                                                                                                                                                                                                   InterPro; IPR000349;
Pfam; PF00695; vMSA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10413;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hepatitis B virus (subtype adw / strain Japan/pJDW233).
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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                     Score 1715;
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21-JUL-1986
21-JUL-1986
15-DEC-1998
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EMBL; J02205; AAA45524.1; -.
                                                                                                                                                                                                                                                                                                                                                                             hepatitis B virus surface antigen."; Nature 280:815-819(1979).
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Valenzuela P., Rutter W.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=79244739; PubMed=471053; Valenzuela P., Gray P., Quiroga
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Viruses; Retroid v
NCBI_TaxID=10408;
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VMSA_HPBV2
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                                                                                                                                               A03706; SAVLVD.
rPro; IPR000349; Hepvir_surfag
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15
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177
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(Rel. 01, Last sequence up
(Rel. 37, Last annotation
ce antigen precursor.
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viruses; Hepadn
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400
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MAJOR SURFACE ANTIGEN.
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
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   CRC64;
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                 (POTENTIAL)
(POTENTIAL)
(POTENTIAL)
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Matches

Conservative

14; Mismatches Score 1713; DB 1; Pred. No. 2.2e-114;

25

Indels

<u>.</u>

Gaps

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Length

400;

Query Match Best Local Similarity

88.6%;

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01-AUG-1990
01-AUG-1992
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                                                                                                                                                                                                                                                                           Okamoto H., Tsuda F., Sakugawa H., Sastrosoewignjo R
Miyakawa Y., Mayumi M.;
"Typing hepatitis B virus by homology in nucleotide
comparison of surface antigen subtypes.";
J. Gen. Virol. 69:2575-2583(1988).
                                    CARBOHYD
CARBOHYD
CARBOHYD
CARBOHYD
SEQUENCE
                                                                                                                            Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                Major surface
                                                                                                                                                 PIR; H28925; SAVLJ2
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Okamoto H., Tsuda F., Sakugawa H.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
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                                                                                                                             PF00695; vMSA;
Similarity
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                                                                                                                                         IPR000349;
                                                                                                                                                                                                                                                                                                                                                                               virus (subtype adw / strain Okinawa/pODW282).
troid viruses; Hepadnaviridae; Orthohepadnavirus
                                                                                                                                                                                                                                                                                                                                                                                                                (Rel. 15, Created)
(Rel. 15, Last sequence up)
(Rel. 23, Last annotation)
ce antigen precursor.
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166
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    88.3%;
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                                     MAJOR SURFACE ANTIGEN.
N-LINKED (GLCNAC. ..)
N-LINKED (GLCNAC. ..)
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N-LINKED (GLCNAC. ..)
CECACODDA3D84A10 CRC6-
    Score 1704; DB 1;
Pred. No. 9.3e-114;
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                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUL-1986 (Rel. 01, Created)
01-AUG-1992 (Rel. 23, Last sequence up
01-NOV-1997 (Rel. 35, Last annotation
                                                                                                                                                                                                                                                                                                              MEDLINE=83168919; PubMed=6300776;
MEDLINE=83168919; PubMed=6300776;
Ono Y., Onda H., Sasada R., Igazashi K., Sugino Y., Nishioka
Ono Y., Onda H., Sasada R., Igazashi K., Sugino Y., Nishioka
                                                                                                                                                                                                                                                                                    "The complete nucleotide sequences of the cloned hepatitis DNA; subtype adr and adw."; Nucleic Acids Res. 11:1747-1757(1983).
                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=106821;
                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis B virus (subtype adw).
                                                                                                                                          InterPro; IPR000349; Hepvir_surfAg.
Pfam; PF00695; vMSA; 1.
                                                                                                                                                                         EMBL; V00866; CAA24233.1; ALT_INIT.
                                                                                                                                                                                                                                                                                                                                                                           Viruses; Retroid
                                                                                                                                                                                                                                                                                                                                                                                                            Major surface
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                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
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           Similarity
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 Conservative
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          87.9%;
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  14;
                                           MAJOR SURFACE ANTIGEN.
N-LINKED (GLCNAC. .)
N-FE9EF2E7EEFC58B5 CRC6
  Pred. No. 4e-113;
Nismatches
                       Score 1695;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LT 12
HPBVI
                                                                                                                                                                                                                                                                                                                                                  MEDLINE=89010694; PubMed=3171552; Okamoto H., Tsuda F., Sakugawa H., Sastrosoewignjo R Miyakawa Y., Mayumi M.; "Typing hepatitis B virus by homology in nucleotide comparison of surface antigen subtypes."; J. Gen. Virol. 69:2575-2583(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VMSA HPBVI
P17397;
                                                                                                                                                                                           interPro; IPR000349; Hepvir_surfAg
Pfam; PF00695; vMSA; 1.
                                                                                                                                                                                                                  EMBL; D00331; -; NOT ANNOTATED CDS.
PIR; I28925; SAVLJ3.
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01-AUG-1990 (Rel. 15, Last sequence update)
01-AUG-1992 (Rel. 23, Last annotation update)
Major surface antigen precursor.
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6 GAGAPGIGFTPPHGGILGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ
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88.0%; Pred. No. 2.4e
tive 12; Mismatches
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N-LINKED (GLCNAC...)

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AC P17101;
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DT 01-AUG-1990 (
DT 01-AUG-1992 (
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                   119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; X51970; CAA36230.1; -. PIR; S10383; SAVLKS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Koechel H.G., Schueler A., Lottmann S., Thomssen R., Submitted (FEB-1990) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis B virus (subtype adw / strain 991).
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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                          AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                          ĠVĠĀŔĠŖĠŔŢŔŖĦĠĠVĹĠŴŚŖŎĀŎĠĸĹŢŖVSŢĮŖŶŔĀŚĀNŖŎŚĠŖŎŖŢŖĬĸŖĸĸĸŎ
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(Rel. 15, Last sequence update)
(Rel. 23, Last annotation updat
ce antigen precursor.
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87.1%;
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N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
N-LINKED (GLCNAC...)
                                                                                                                                                                                    Score 1684; DB 1;
Pred. No. 2.5e-112;
6; Mismatches 28;
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Matches 289; Conserv
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                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use by non-profit institutions as long as its content is in no war modified and this statement is not removed. Usage by and for commercia entities requires a license agreement (See http://www.isb-sib.ch/announce or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                   Antigen.
PROPEP
CHAIN
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Viruses, Retroid viruses, Hepadnaviridae, Orthohepadnavirus.
                                                                                                                                                                                                                                                                                                                      EMBL; X69798; CAA49455.1; -.
InterPro; IPR000349; Hepvir
Pfam; PF00695; vMSA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=45410;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Identification of a new hepatitis B virus (HBV) genotype from that expresses HBV surface antigen subtype adw4.";
J. Gen. Virol. 74:1627-1632(1993).
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Naumann H., Schaefer S., Yoshida C.F.T., Gaspar A.M.C., Repp
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CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQQMLPVCPLIPGSSTTSTGPCRTC
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MEDLINE-88297159; PubMed=2841200;

MILLINE-88297159; PubMed=284120
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A; Residues: 264-343 «NOR» A; Experimental source: subty C; Genetics: pre-S1/pre-S2/S A; Gene: pre-S1/pre-S2/S C; Superfamily: hepatitis B v C; Keywords: glycoprotein; su: F;109-389/Product: major su: F;164-389/Product: major su: F;44,112,166/Binding site: ca: Query Match Best Local Similarity 100 Matches 342; Conservative Oy 6 GAGAFGLGFTPPHGGLLGG MATCHES 342; CONSTTFHQTLODPRVI Db 48 GAGAFGLGFTPHTLJODPRVI OY 66 AMOMNSTTFHQTLODPRVI Db 108 AMQMNSTTFHQTLODPRVI OY 126 TSGFLGPLLVLQAGFFLLT Db 168 TSGFLGPLLVLQAGFFLLT Db 168 TSGFLGPLLVLQAGFFLLT Db 168 TSGFLGPLLVLQAGFFLLT Db 168 TSGFLGPLLVLQAGFFLLT Db 228 CPPTCPGYRWMCLRRFIIF	A;Residues: 1-389 <bic> A;Cross-references: GB:X0249 R;Norder, H.; Courouce, A.M. J. Gen. Virol. 73, 3141-3145 A;Title: Molecular basis of A;Reference number: PQ0453; A;Accession: PQ0585 A;Molecule type: DNA A;Molecule type: DNA</bic>	RESULT 1 Large surface antigen - hepa large surface antigen - hepa large surface antigen - hepa N;Contains: major surface ar C;Species: hepatitis B virus A;Note: host Homo sapiens (u C;Date: 17-Mar-1987 #sequenc C;Accession: A03704; PQ0585 R;Bichko, V.; Pushko, P.; Dr FEBS Lett. 185, 208-212, 198 A;Title: Subtype ayw variant A;Reference number: A05237; A;Accession: A03704 A;Molecule type: DNA A;Molecule type: DNA		1256 65.1 1253 65.0 1253 65.0 1250 64.8 1250 64.8 1249 64.7	30 1269 65.8 22 31 1267 65.7 22 32 1267 65.6 22 33 1265 65.6 22 34 1264 65.5 22 35 1264 65.5 22 36 1263 65.5 22 37 1269 65.3 22 38 1259 65.3 22

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ALIGNMENTS

98.5%; Score 1901; DB 1; Length 389; 100.0%; Pred. No. 3.3e-128; ive 0; Mismatches 0; Indels 0; virus surface antigen surface antigen surface antigen (gene pre-S2/S) #status predicted <DSA> urface antigen (gene S) #status predicted <MSA> carbohydrate (Asn) (covalent) #status predicted 496; NID:g62280; PIDN:CAB41701.1; PID:g4704321 W.; Magnius, L.O. 15, 1992 E hepatitis B virus serotype variations within the four major; MUID:93107848; PMID:1469353 patitis B virus (subtype ayw, strain pHB320) antigen; middle surface antigen us, HBV IFLFILLLCLIFLLVLLDYQGMLPVCPLIFGSSTTSTGPCRTC 245 GWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65 Dreilina, D.; Pumpen, P.; Gren, B. 985 nce_revision 17-Mar-1987 #text_change 28-Jul-2000 TRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS JTRILTIPOSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185 ype ayw2, Tav of hepatitis B virus: DNA primary structure analysis. MUID:85204397; PMID:3996597 Gaps 227 0,

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surface antigen - hepatitis B virus (su N;Alternate names: HBs antigen N;Contains: surface antigen pre-Si (lar. C;Species: hepatitis B virus, HBV A;Variety: subtype ayw, patient C C;Date: 20-Feb-1995 #sequence_revision
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A;Residues: 1-389 <LAI>
A;Cross-references: EMBL:X65257; NID:959429; PIDN:CAA46349.1;
A;Experimental source: subtype ayw, patient C
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Accession: $20745

R;Lai, M.E.; Mazzoleni, A.P.; Balestrieri, A.; Melis, A.; Porru, 
submitted to the EMBL Data Library, March 1992

A;Description: Sequence analysis of HBV genomes isolated from pa 
A;Reference number: $20745

A;Accession: $20745
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large surface antigen - hepatitis B virus (subtype ayw, isolate N,Contains: major surface antigen; middle surface antigen C,Species: hepatitis B virus, HBV C,Species: subtype ayw, isolate patient C1005 A,Variety: subtype ayw, isolate patient C1005 C;Date: 20-reb-1995 #sequence_revision 06-Dec-1996 #text_change
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Matches 338
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;1-108/Domain: pre-52 domain #status predicted <PRE2>
;109-389/Product: surface antigen pre-52 (middle envelope protein) #s;109-163/Domain: pre-52 domain #status predicted <PRE2>
;164-389/Product: surface antigen S (small envelope protein) #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Keywords: surface antigen
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Pred. No. 6.3e
2; Mismatches
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C;Accession: S32202
R;Preisler-Adams, S.; Schlayer, M.J.; Peters, T.; Hettler, F.; Gerok, W.; Rasenack, J. submitted to the EMBL Data Library, March 1993
A;Description: Identification and sequence analysis of hepatitis B virus DNA in immuno A;Reference number: S32202
A;Accession: S32202
A;Accession: S32202
A;Accession: S32202
A;Rolecule type: DNA
A;Residues: 1-389 < PRE>
A;Cross-references: EMBL:X72702
A;Cross-references: EMBL:X72702
A;Experimental source: subtype ayw, isolate patient C1005
C;Genetics:
A;Gene: pre-S1/pre-S2/S
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: surface antigen
F;1-389/Product: major surface antigen (gene pre-S2/S) #status predicted <DSA>
F;109-389/Product: major surface antigen (gene pre-S2/S) #status predicted <MSA>
F;164-389/Product: major surface antigen (gene S) #status predicted <MSA>
                                                                                                                                                                                                                                                                                                                                   Burface antigen - hepatitis B virus (subtype ayw4)
N,Alternate names: envelope protein; HBs antigen
N;Contains: surface antigen pre-Si (large envelope
C;Species: hepatitis B virus, HBV
A;Variety: subtype ayw4
C;Date: 23-Nov-1994 #sequence_revision 12-May-1995
C;Accession: $47407; JQ2071
R;Plucienniczak, A.
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                                 A;Molecule type: DNA
A;Residues: 1-389 <PUV
A;Residues: 1-389 <PUV
A;Cross-references: EMBL:235716; NID:g527435; PIDN:CAA84788.1; PID:g527439
A;Cross-references: Subtype ayw4
A;Experimental source: subtype ayw4
R;Norder, H; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, R;Norder, H; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, J., Gen. Virol. 74, 1341-1348, 1993
A;Title: Genetic relatedness of hepatitis B viral strains of diverse geogr.
A;Reference number: JQ2044; MUID:93329382; PMID:8336122
A;Reference subser: JQ2044; MUID:93329382; PMID:8336122
A;Accession: JQ2071
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A; Molecule type: DNJ
A; Residues: 164-389
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337; Conserv
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98.5%;
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Pred. No. 1.2e-126;
2; Mismatches 3;
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d sequencing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein); surface
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ANOR.

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A;Molecule type: DNA
A;Residues: 1-389 <GAL>
A;Cross-references: GB:J02203
R;Porder, H.; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahv
J. Gen. Virol. 74, 1341-1348, 1993
A;Title: Genetic relatedness of hepatitis B viral strains of diverse of A;Reference number: J02044; MUID:93329382; PMID:8336122
A;Accession: J02064
A;Molecule type: DNA
A;Residues: 164-389 <NOR>
A;Accessioues: 164-389 <NOR>
A;Residues: 164-389 <NOR>
A;Title: Molecular basis of hepatitis B virus serotype variations with A;Residues: 164-389 <NORS
A;Residues: 164-389 <NORS
A;Residues: 264-343 <NOS>
A;Residues: 264-343 <NOS>
A;Residues: 264-343 <NOS>
A;Residues: 264-343 <NOS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   large surface antigen - hepatitis B virus (subtype ayw)
N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBC
C;Date: 18-Dec-1981 #sequence revision 18-Dec-1981 #text_change 07-May-1999
C;Accession: A03703; JQ2064; FQ0591
R;Galibert, F:; Mandart, B:; Fitoussi, F:; Tiollais, P:; Charnay, P.
Nature 281, 646-650, 1979
A;Title: Nucleotide sequence of the hepatitis B virus genome (subtype ayw) i
A;Reference number: A93214; MUID:81012091; PMID:399327
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C;Superfamily: hepatitis B virus surface antigen
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: surface antigen
F;1-389/Product: surface antigen pre-S1 (large envelope protein) #status predicted <PS
F;1-108/Domain: pre-S1 domain #status predicted <PRE1>
F;109-389/Product: surface antigen pre-S2 (middle envelope protein) #status predicted
F;109-163/Domain: pre-S2 domain #status predicted <PRE2>
F;109-163/Domain: pre-S2 domain #status predicted <PRE2>
F;109-163/Domain: pre-S2 domain #status predicted <SAG>
F;109-163/Domain: pre-S2 domain #status predicted <PRE3-
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   A; Experimental source:
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98.2%;
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348, 1993
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N;Contains: major Burface antigen, middle surface antigen C;Species: hepatitis B virus, HBV C;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change : C;Accession: B34773; JQ2082 R;Tong, S.; Li, J.; Vitvitski, L.; Trepo, C. Virology 176, 596-603, 1990 A;Title: Active hepatitis B virus replication in the presence of A;Reference number: A34773; MUID:90266476; PMID:2345966 A;Accession: B34773
A;Status: translation not shown
                                                G;Keywords: glycoprotein; surface antigen F;109-389/Product: middle surface antigen (gene pre-S2/S) #status predicted F;164-389/Product: major surface antigen (gene S) #status predicted <MSA>F;4,112,166/Binding site: carbohydrate (Asn) (covalent) #status predicted
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A.Residues: 1-389 <TON>
A;Cross-references: EMBL:M32138; NID:g329667; PIDN:AAA45502.1; PID:g329669
A;Cross-references: EMBL:M32138; NID:g329667; PIDN:AAA45502.1; PID:g329669
R.Morder, H.; Hammas, B.; Lee, S.D.; Bile, K.; Courouce, A.M.; Mushahwar, I.K.; Magnius, J. Gen. Virol. 74, 1341-1348, 1993
J. Gen. Virol. 74, 1341-1348, 1993
A;Fitle: Genetic relatedness of hepatitis B viral strains of diverse geographical origin A;Reference number: JQ2044; MUID:93329382; PMID:8336122
A,Accession: JQ2082
A,Accession: JQ2082
                                                                                                                                                                       A;Gene: pre-S1/pre-S2/S
A;Introns: 111/3
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A; Residues: 164-389 < NOR>
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C,Superfamily: hepatitits B virus surface antigen
C;Keywords: glycoprotein; surface antigen
C;Keywords: glycoprotein; surface antigen
F;109-389/Product: middle surface antigen (gene pre-S2/S) #status predicted
F;164-389/Product: major surface antigen (gene S) #status predicted <MSA>
F;164-389/Product: major surface antigen (gene S) #status predicted
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Pred. No. 7.3e~126;
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Best Local Query Match

Similarity

Conservative

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Mismatches

.3e-126; DB 1;

Indels

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Gaps

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Length 389;

Score 1868; Pred. No. 7.

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C;Accession: $20753
R;LAi, M.E.; Mazzoleni, A.P.; Balestrieri, A.; Melis, A.; Porru, Bubmitted to the EMBL Data Library, March 1992
A;Description: Sequence analysis of HBV genomes isolated from pat A;Reference number: $20745
A;Accession: $20753
A;Accession: $20753
A;Molecule type: DNA
A;Residues: 1-389 <LAI>
A;Cross-references: EMBL:X65259; NID:g59439; PIDN:CAA46357.1; PID
A;Experimental source: subtype ayw, patient E
C;Genetics:
A;Gene: $
C;Genetics:
A;Gene: $
C;Superfamily: hepatitis B virus surface antigen
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: surface antigen pre-S1 (large envelope protein):
F;1-189/Domain: pre-S1 domain #status predicted <PRE1>
F;109-163/Jomain: pre-S2 domain #status predicted <PRE2>
F;109-189/Product: surface antigen S (small envelope protein) #st.
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N;Alternate names: envelope protein; HBs antigen
N;Contains: surface antigen pre-S1 (large envelope protein);
C;Species: hepatitis B virus, HBV
A;Variety: subtype ayw, patient E
C;Pate: 20-Feb-1995 #sequence_revision 01-Nov-1996 #text_chan
C;Accession: $20753
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;1-389/product: surface antigen pre-S1 (large envelope protein) #81
;1-389/product: surface antigen pre-S1 (middle envelope protein)
;109-389/product: surface antigen pre-S2 (middle envelope protein)
;109-163/Domain: pre-S2 domain #status predicted <PRE2>
;164-389/Product: surface antigen S (small envelope protein) #statu
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Best Local
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                       CPPTCPGYRWMCLRRPIIFLFILLLCLIFLLVLLDYQGMLFVCFLIFGSSTTSTGPCRTC
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97.71;
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Pred. No. 2.
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2.3e-125;
1es 5;
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surface antigen - hepatitis B virus (subt N;Alternate names: envelope protein; HBs N;Contains: surface antigen pre-Si (large C;Species: hepatitis B virus, HBV A;Variety: subtype ayw, patient P C;Date: 06-Jan-1995 #Bequence_revision 06 C;Accession: S41871 R;Lai, M.E.; Mazzoleni, A.P.; Balestrieri submitted to the EMBL Data Library, Janua A;Description: Sequence analysis of HBV g

06-Jan-1995

#text_change

26-Aug-1999

B virus (subtype ayw, patient protein; HBs antigen pre-Sl (large envelope protei

protein);

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A.P., Balestrieri,

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large surface antigen - hepatitis B virus (subtype adyw)
large surface antigen; middle surface antigen
N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBV
C;Species: hepatitis B virus, HBV
C;Accession: A93217; A03703
C;Accession: A93217; A03703
R;Pasek, M.; Goto, T.; Gilbert, W.; Zink, B.; Schaller, H.; MacKsy, P.; Leadbett
Nature 282, 575-579, 1979
A;Title: Hepatitis B virus genes and their expression in E. coli.
A;Reference number: A93217; MUID:81012115; PMID:399329
A;Accession: A93217
A;Accession: A93217
A;Coss-references: GB:J02202
C;Genetics:
B.Gonetics:
C.Genetics:
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C;Superfamily: hepatitis B virus surface antigen
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: glycoprotein; surface antigen
F;109-389/Product: middle surface antigen (gene pre-S2/S) #status predicted
F;104-389/Product: major surface antigen (gene S) #status predicted <MSA>
F;164-389/Product: major surface antigen (gene S) #status predicted cMSA>
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GLSPIVMLSVIWMMYWGPSLYSILSPFLPLLPIFFCLWAYI
                        GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI
                                                               TTPAQGISMYPSCCCTKPSDGNCTCIPIPSSMAFGKFLMEMASARFSWLSLLVPFVQWFV
                                                                                  TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVOWFV
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97.1%;
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Pred. No. 1.2e-124;
2; Mismatches 8;
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A; Description: Sequence analysis of HBV genomes 1, A; Reference number: $41869
A; Accession: $41869
A; Molecule type: DNA
A; Residues: 1-378 < LAIS
A; Cross-references: EMBL: X77309; NID: 9452611; PNI
A; Cross-references: EMBL: X77309; NID: 9452611; PNI
A; Experimental source: subtype ayw, patient A
C; Genetics:
A; Genetics:
C; Superfamily: hepatitis B virus surface antigen
C; Keywords: surface antigen
F; 1-378/Product: surface antigen pre-$1 (large e)
F; 1-108/Domain: pre-$1 domain #status predicted.
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A;Accession: S41871
A;Molecule type: DNA
A;Residues: 1-389 < LAI>
A;Residues: 1-389 < LAI>
A;Experimental source: Subtype ayw, patient p
C;Genetics:
A;Gene; S
C;Superfamily: hepatitis B virus surface antigen
C;Keywords: surface antigen
F;1-389/Product: Surface antigen pre-S1 (large en
F;1-108/Domain: pre-S1 domain #status predicted
F;109-163/Domain: pre-S2 domain #status predicted
F;109-163/Domain: pre-S2 domain #status predicted
                    F;1-378/Product: surface antigen pre-S1 (large envelope protein)
F;1-108/Domain: pre-S1 domain #status predicted <PRE1>
F;109-378/Product: surface antigen pre-S2 (middle envelope protein)
F;109-163/Domain: pre-S2 domain #status predicted <PRE2>
                                                                                                                                                                                                                                                                                                                                                    N;Contains: surface antigen pre-S1 (lat
C;Species: hepatitis B virus, HBV
A;Variety: subtype ayw, patient A
C;Date: 06-Jan-1995 #sequence_revision
C;Accession: S41869
R;Lai, M.E.; Mazzoleni, A.P.; Balestrie
                                                                                                                                                                                                                                                                                                        submitted to the EMBL Data Library,
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;Superfamily: hepatitis B virus surrace antigen
;Kupwords: surface antigen
;kaywords: surface antigen pre-S1 (large envelope protein) #status
?;1-189/Product: surface antigen pre-S2 (middle envelope protein) #stat
P;109-389/Product: surface antigen pre-S2 (middle envelope protein) #stat
F;109-163/Domain: pre-S2 domain #status predicted <PRE2>
F;109-163/Domain: surface antigen S (small envelope protein) #status pre-
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Lai M.E.; Mazzoleni, A.P.; Balestrieri, A.
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Pred. No. 3.1e-121
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A;Cross-references: EmBL:X65258; NID:g59434; PII
A;Experimental source: subtype ayw, patient CI
A;Note: typical methionine for the beginning of C;Genetics:
A;Gene: S
C;Superfamily: hepatitis B virus surface antige C;Ecywords: surface antigen C;Ecywords: surface antigen pre-S1 (large F;1-389/Product: surface antigen S (small env
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A;Variety: subtype ayw, patient CI
C;Date: 20-Feb-1995 #sequence_revision 01-Nov-1996 #text_change
C;Accession: S20749
R;Lai, M.E.; Mazzoleni, A.P.; Balestrieri, A.; Melis, A.; Porru,
                                                                                                                                                                                                                                                                                                                                                                                                         A; Residues: 1-389 < LAI>
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A;Accession: S20749
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SCPPTCPGYRMMCLRRFIIFLFILLLCLIFLLVLDYQGMLPVCPLIFGSSTTSTGPCRT
                                          ITSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFRGGTTVCLGQNSQSPTSNHSPT
                                                                                                                  QAMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMEN
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Pred. No. 8.4e-121;
tl; Mismatches 13;
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Pred. No. 5.9e-121;
3; Mismatches 4;
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large surface antigen - hepatitis B virus (subtype adr)
N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBV
C;Species: hepatitis B virus, HBV
C;Date: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 07-May-1999
C;Date: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 07-May-1999
C;Accession: A03705; S04569; J02107; P00508
R;Onc, Y; Onda, H; Sasada, R; Jgarashi, K; Sugino, Y; Nishioka, K.
N; Onco, Y; Onda, H; Sasada, R; Jgarashi, K; Sugino, Y; Nishioka, K.
N; Nishioka, K; Nishioka, Nishioka
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A;Title: Genetic relatedness of hepatitis B viral strains of diverse geograph a; Reference number: JQ2044; MUID:93329382; PMID:8336122
A; Accession: JQ2107
A; Molecule type: DNA
A; Residues: 175-400 < NOR>
A; Residues: 175-400 < NOR>
A; Residues: 175-400 < NOR>
A; Resperimental source: genogroup C, subtype adr, strain pBRHBadr4
A; Experimental source: genogroup C, subtype adr, strain pBRHBadr4
A; Reference number: P00453; MUID:93107848; PMID:1469353
A; Reference number: P00453; MUID:93107848; PMID:1469353
A; Accession: P00653; MUID:93107848; PMID:1469353
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R;Rho, H.M.; Kim, K.; Hyun, S.W.; Kim, Y.S.
Nucleic Acids Res. 17, 2124, 1989
A;Title: The nucleotide sequence and reading frames of a
A;Reference number: S04568; MUID:89183619; PMID:2928116
A;Accession: S04569
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A; Residues: 275-354 <NO2>
A; Experimental
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A;Residues: 1-50,'K',52-66,'YP',69-129,'V',131-142,'P',144-164,'S',166-176,'S',178-338,
A;Cross-references: EMBL:X14193
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Best Local Sim
Matches 315;
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;Gene: pre-S1/pre-S2/S
;Superfamily: hepatitis B virus surface antigen
;Superfamily: hepatitis B virus surface antigen
;Reywords: glycoprotein; surface antigen (gene pre-S2/S) #status predicted
;1120-400/Product: middle surface antigen (gene S) #status predicted <MSA>
;115-400/Product: major surface antigen (gene S) #status predicted
;15,123,177/Binding site: carbohydrate (Asn) (covalent) #status predicted
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                                                                                                                                                                                                                                                                                GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
TSGFLGPLLVLQAGEFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                                                     AMOWNSTTFHOALLDPRVRGLYFPAGGSSSGTVNPVPTTASPISSIFSRTGDPAPNMENT 178
                                                                                                                                               AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                                                                                                                                             h 91.8%; Score 1771; DB 1; Length 400;
Similarity 92.1%; Pred. No. 6.1e-119;
15; Conservative 10; Mismatches 17; Indels
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large surface antigen - hepatitis B virus
N;Contains: major surface antigen; middle surface antigen
N;Contains: major surface antigen; middle surface antigen
C;Species: hepatitis B virus, HBV
C;Species: hepatitis B virus, HBV
C;Species: no.7un-1989 #sequence_revision 30-Jun-1989 #text_change 16-Jun-2000
C;Date: 30-Jun-1989 #sequence_revision 30-Jun-1989 #text_change of the genome of a hepatitis B virus isolated if
A;Coession: C28885
A;Coession: C28885
A;Coession: A92796; MUID:88258473; pMID:2838576
A;Courouce, AM:; Mushahwar, I.K.; Magnius,
A;Rociesion: J91
A;Coession: J91
A;Coession: J92
A;Coession: J93
A;Coession: J92
A;Coession: J93
A;Coes
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C;Superfamily: hepatitis B virus surface antigen
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                                                                                                                CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCKTC
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                                TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
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Pred. No. 8.2e-118;
9; Mismatches 20;
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GLSPTVWLSVIWMWYWGPSLYSILSPFLPLLPIFFCLWVYI 347

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surface antigen - hepatitis B virus (subtype ayw, p N,Alternate names: envelope protein; HBs antigen N;Contains: surface antigen pre-S1 (large envelope C;Species: hepatitis B virus, HBs A;Variety: subtype ayw, patient M C;Date: 06-Jan-1995 #sequence_revision 01-Nov-1996 C;Accession: S41870
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A; Introns: 165/2
C; Superfamily: he
C; Keywords: surfa
F; 1-400/Product:
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A;Title: The complete nucleotide sequence of hepatitis B virus, subtype A;Reference number: $35527; MUID:93096607; PMID:1461746
A;Recession: $35528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: EMBL:D12980; NID:9221500; PIDN:BAA02359.1; PID:9221502
A;Experimental source: subtype adr
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, Au;C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Status: nucleic ac:
A;Molecule type: DNA
A;Residues: 1-400 <M
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C;Date: 09-Dec-1993 #sequence_revision 10-Nov-1995 #text_change 20-Jun-2000
C;Accession: S35528
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Keywords: surface antigen
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C;Superfamily: hepatitis B virus surface antigen
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A;Experimental source: subtype ayw, patient M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           submitted to the EMBL Data Librar A;Description: Sequence analysis A;Reference number: $41869 A;Accession: $41870
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A; Residues: 1-378 <LAI>
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Best Local Similarity
Matches 311; Conserv
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;1-108/Domain: pre-S1 do
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Pred. No. 2.5e-116;
7; Mismatches 13;
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HBV genomes isolated from
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               Score
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Gapop 10.0 , Gapext 0.
      404799 seqs, 80919614 residues
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     DB
          PCT-US02-32263-46
US-10-287-994-46
US-10-287-994-19
US-10-326-908-19
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US-10-326-908-19
US-10-326-908-20
US-09-863-054-22
US-09-724-676-50956
US-09-724-6768-50956
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US-09-724-6768-50959
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US-10-311-034-15
US-10-311-034-15
US-10-284-130-13
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                                                                                                                         Sequence 46, Appl
Sequence 21, Appl
Sequence 21, Appl
Sequence 19, Appl
Sequence 18, Appl
Sequence 22, Appl
Sequence 22, Appl
Sequence 50955, A
Sequence 50957, A
Sequence 50957, A
Sequence 50957, A
Sequence 50957, A
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Sequence 28, Appl
Sequence 327, Appli
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		Sequence 86526, A	Sequence 86525, A	Sequence 86515, A					Sequence 86515. A	Sequence 86514, A	sequence 86513, A	, 71000	24	Sequence 30. Appl	Sequence 3, Appli	sequence 3355, Ap		ر د د	Sequence 4. Appli	rdderice ac, Appr	9 6	Seguence An appl	

ALIGNMENTS

APPLICANT: Bayer, Robert APPLICANT: Bayer, Robert APPLICANT: Hakes, Daryn APPLICANT: Chen, Xi ITITE OF INVENTION: REMODELING AND GLYCOCONJUGATION OF PEPTIDES FILE REFERENCE: 040853-01-505000 CURRENT PILING DATE: 2002-10-31 PRIOR APPLICATION NUMBER: US 60/328,523 PRIOR APPLICATION NUMBER: US 60/344,692 PRIOR APPLICATION NUMBER: US 60/344,692 PRIOR FILING DATE: 2001-10-10 PRIOR APPLICATION NUMBER: US 60/334,233 PRIOR FILING DATE: 2001-11-08 PRIOR APPLICATION NUMBER: US 60/334,301 PRIOR APPLICATION NUMBER: US 60/334,301 PRIOR APPLICATION NUMBER: US 60/334,301 PRIOR APPLICATION NUMBER: US 60/387,292 PRIOR APPLICATION NUMBER: US 60/387,292 PRIOR FILING DATE: 2002-06-07 PRIOR APPLICATION NUMBER: US 60/391,777 PRIOR APPLICATION NUMBER: US 60/396,594 PRIOR APPLICATION NUMBER: US 60/396,594 PRIOR APPLICATION NUMBER: US 60/396,594 PRIOR APPLICATION NUMBER: US 60/404,249 PRIOR FILING DATE: 2002-08-16 PRIOR APPLICATION NUMBER: US 60/404,249 PRIOR FILING DATE: 2002-08-16 PRIOR APPLICATION NUMBER: US 60/404,249 PRIOR FILING DATE: 2002-08-28 NUMBER OF SEQ ID NOS: 62 SOFTWARE: Patentin version 3.1 LENGTH: 406 LENGTH: 406 LENGTH: 400 ; TYPE: PRT ; ORGANISM: Hepatitis PCT-US02-32263-46 RESULT 1 PCT-US02-32263-46 Sequence 46, Application PC/TUS0232263 GENERAL INFORMATION: APPLICANT: Neose Technologies, Inc. APPLICANT: DeFrees, Shawn APPLICANT: Zopf, David B virus

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h 91.0%; Score 1756; DB 1; Similarity 91.2%; Pred. No. 4.5e-106; 12; Conservative 11; Mismatches 19;

Length 400; Indels

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HAPPLICANI: CHEM, A1
TITLE OF INVENTION. REMODELING AND GLYCOCONJUGATION OF PEPTIDES
PILE REFERENCE: 040853-01-5052-00
PILE REFERENCE: 040853-01-5052-00
CURRENT APPLICATION NUMBER: US/10/287,994
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US 60/328,523
PRIOR APPLICATION NUMBER: US 60/344,692
PRIOR APPLICATION NUMBER: US 60/344,692
PRIOR APPLICATION NUMBER: US 60/387,292
PRIOR APPLICATION NUMBER: US 60/387,292
PRIOR APPLICATION NUMBER: US 60/391,777
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PRIOR APPLICATION NUMBER: US 60/396,594
PRIOR APPLICATION NUMBER: US 60/396,594
PRIOR FILING DATE: 2002-06-25
PRIOR FILING DATE: 2002-08-25
PRIOR APPLICATION NUMBER: US 60/404,249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               306
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                                126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTPAQGTSMYDSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TSGPLGPLLVLQAGPFLLTRILTIPQSLDSWTSLNPLGGAPTCPGQNSQSPTSNHSPTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GLSPTVWLSVIWMMWYWGPSLYSILSPELPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TIPAQCISMEPSCCCTKPSDGNCTCIPIPSSWAFARFLWEWASVRESWLSLLVPFVQWFA 358
                                                                                                                                                                                                       GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRWTHPQ 65
                                                                                                                                 AMQWNSTTFHQTLQDPRVRGLYPPAGGSSSGTVNPVPTTVSPISSIPSRIGDPALMMENI 125
TSGFLGPLLVLQAGPFLLTRILTIPQSLDSWMTSLNFLGGAPTCPGQNSQSPTNNHSPTS
                                   T8GFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                                                   Bayer, Robert
Bowe, Caryn
Hakes, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hakes, Da
Chen, X1
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                                                                                                                                                                                                                                                                                                        91.0%;
iilarity 91.2%;
Conservative 11
                                                                                                                                                                                                                                                                                                           ; Score 1756; DB 6;
; Pred. No. 4.5e-106;
11, Mismatches 19;
                                                                                                                                                                                                                                                                                                                                                                    Length 400;
                                                                                                                                                                                                                                                                                                                      Indels
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             238
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TITLE OF INVENTION: Designing Immunogens
FILE REFERENCE: 117-433 / N78027D
CURRENT APPLICATION NUMBER: US/10/326,908
CURRENT FILING DATE: 2002-12-33
PRIOR APPLICATION NUMBER: US 10/110,036
PRIOR APPLICATION NUMBER: US 10/110,036
PRIOR APPLICATION NUMBER: US 10/110,036
PRIOR APPLICATION NUMBER: GB 0007789.1
PRIOR APPLICATION NUMBER: GB 0007789.1
PRIOR APPLICATION NUMBER: GB 9923902.2
PRIOR APPLICATION NUMBER: GB 9923902.2
PRIOR FILING DATE: 2000-03-30
PRIOR FILING DATE: 1999-10-08
NUMBER OF SEQ ID NO 21
LENGTH: G7
TYGE: DBT
TYGE: D
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GENERAL INFORMATION:
APPLICANT: Christopher JONES
APPLICANT: Andrew BACON
APPLICANT: Gill DOUCE
APPLICANT: Mark PAGE
APPLICANT: Mark PAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              먕
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APPLICANT: Christopher JONES
APPLICANT: Andrew BACON
APPLICANT: Andrew BACON
APPLICANT: Gill DOUCE
APPLICANT: Mark PAGE
TITLE OF INVENTION: Designing Immunogens
FILE REFERENCE: 117-433 / N78027D
FILE REFERENCE: 117-433 / N78027D
CURRENT APPLICATION NUMBER: US10/326,908
CURRENT APPLICATION NUMBER: US10/110,036
PRIOR APPLICATION NUMBER: US10/110,036
PRIOR APPLICATION NUMBER: CCT/GB00/03857
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: GB 0007789.1
PRIOR APPLICATION NUMBER: GB 0007789.2
PRIOR APPLICATION NUMBER: GB 9923902.2
PRIOR APPLICATION NUMBER: GB 9923902.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 15.6%; Score 300; DB 6; 1
Best Local Similarity 100.0%; Pred. No. 8.3e-13;
Matches 57; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   246
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TIPAQGTSMEPSCCCTKESDGNCTCIPIPSSWAFARFLWEWASVRESWLSLLVPFVQWFA 358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CPPICPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLLPGTSTTSTGPCKTC
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APPLICANT: Christopher JONES
APPLICANT: Andrew BACON
APPLICANT: Andrew BACON
APPLICANT: Gill DOUCE
APPLICANT: Mark PAGE
TITLE OF INVENTION: Designing Immunogens
FILE REFERENCE: 117-433 / NT8027b
CURRENT APPLICATION NUMBER: US/10/326,908
CURRENT FILING DATE: 2002-12-23
PRIOR APPLICATION NUMBER: US 10/110,036
PRIOR FILING DATE: 2002-04-08
PRIOR APPLICATION NUMBER: PCT/GB00/03857
                                                                                                                                                                                                                              RESULT 6
US-10-326-908-20
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LENGTH: 63
                                                                                                                                                                                               Sequence 20, Application US/10326908 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-326-908-19
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Best Local Similarity
Matches 22; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18, Application US/10
GENERAL INFORMATION:
APPLICANT: Christopher JONES
APPLICANT: Andrew BACON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 10/110,036
PRIOR EILING DATE: 2002-04-08
PRIOR APPLICATION NUMBER: PCT/GB00/03857
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: GB 0007789.1
PRIOR FILING DATE: 2000-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 30
SOFTWARE: MS WORD
SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: GB 9923902.2 PRIOR FILING DATE: 1999-10-08 NUMBER OF SEQ ID NOS: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Designing Immunogens
FILE REFERENCE: 117-433 / N78027D
CURRENT APPLICATION NUMBER: US/10/326,908
CURRENT FILING DATE: 2002-12-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: synthetic peptide immunogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: synthetic peptide immunogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 64
                                                                                                                                                                                                                                                                                              40 GAGAFGLGFTPPHGGLLGWSPQSE 63
                                                                                                                                                                                                                                                                                                                     6 GAGAFGLGFTPPHGGLLGWSPQAQ 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65 QAMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPI88IFSRIGDPALNMB 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
les 57; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gill DOUCE
Mark PAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/10326908
                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                               6.8%; Score 131; DB 6
91.7%; Pred. No. 0.064;

 Mismatches

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Pred. No. 1.le-12;
"" omatches 2;
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US-09-863-054-22
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US-09-863-054-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 20
LENGTH: 68
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: synthetic peptide immunogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                   TELEPHONE: (415) 576-02
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 22;
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: GB 0007789.1
PRIOR FILING DATE: 2000-03-30
PRIOR APPLICATION NUMBER: GB 9923902.2
PRIOR FILING DATE: 1999-10-08
NUMBER OF SEQ ID NOS: 30
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                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 07/935,898
APPLICATION NUMBER: US 08/024,120
APPLICATION NUMBER: US 08/024,120
APPLICATION NUMBER: US 08/396,283
APPLICATION NUMBER: US 08/396,283
FILING DATE: 27-PBB-1995
APPLICATION NUMBER: US 08/463,486
PILING DATE: 05-JUN-1995
APPLICATION NUMBER: US 08/463,486
APPLICATION NUMBER: US 08/463,486
APPLICATION NUMBER: US 08/463,486
APPLICATION NUMBER: US 08/463,486
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nes 22; Conservative
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             TOPOLOGY: linear MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: 5
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                                                                                                                                                                                                                 NAME: Lockyer, Jean M.
REGISTRATION NUMBER: 44,879
REFERENCE/DOCKET NUMBER: 014
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/863,054
FILING DATE: 21-May-2001
CLASSIFICATION: -Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Chisari, Francis V. TITLE OF INVENTION: Peptides f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                          STRANDEDNESS: <Unknown>
                                                                                           TYPE: amino acid
                                                                                                                 LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Francisco
                                                                                                                                                                                                  (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptides for Inducing Cytotoxic T Lymphocyte Responses to Hepatitis B Virus
             SEQ ID NO: 22:
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Sequence 50955, Application US/09724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative spl
TITLE REPERENCE: 129181.4 Compugen
FILE REPERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT ETLING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOOTWARE: Patentin version 3.2
SEQ ID NO 50955
LENGTH: 276
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US-09-724-676-50956
; Sequence 50956, Application US/09724676
; GENERAL IMPORMATION:
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                                                                                                                                                                                   SOFTWARE: PatentIn version 3.2 SEQ ID NO 50956
                                                                            Query Match
Best Local (
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TITLE OF INVENTION: Variants of alternative
FILE REPERENCE: 129181.4 Compugen
CURRENT EDILICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                       TYPE: PRT
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                                                                                                                                                                      LENGTH: 276
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                                                                              Local Similarity
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GGGGGGSFGGGGPPGLGGLFQAGMPKLRSTANRDNDSGGSRP-----PLLPPGGR 82
                                GRGDGAGAFGLGFTPPHGGLL-----
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26.0%; Pred. No. 1
                                                               5.4%; Score 105; DE 26.0%; Pred. No. 12; tive 11; Mismatches
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                                  -----GWSPQAQGILETLPANPP--- 40
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RESULT 10
US-09-724-676-50957
US-09-724-676-50957
Sequence 50957, Application US/09724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129.81.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT APPLICATION STATE: 2000-11-28
UMMBER OF SEQ ID NOS: 97222
SEQ ID NO 50957
LENGTH: 276
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TITLE OF INVENTION: Variants of alternative sp
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
UMMBER OF SEQ ID NOS: 97222
SOFTWARE: Patentin version 3.2
SEQ ID NO 50955
LENGTH: 276
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Best Local Similarity
Matches 57; Conserv
                                                                                                                                                                                                                                                                                                Sequence 50955, Application US/09724676A GENERAL INFORMATION:
                                   Matches
                                                  Best Local
                                                               Query Match
                                                                                                                                   TYPE: PRT
                                                                                                                     ORGANISM: Homo sapiens
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   2 GRGDGAGAFGLGFTPPHGGLL--
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                                   57;
                                                  Similarity
                             5.4%; initarity 26.0%; l
Conservative 11
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Pred. No. 12;
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        ---GWSPQAQGILETLPANPP---
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RESULT 13

US-09-724-676A-50957

i Sequence 50957, Application US/09724676A

i GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative spl

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: US/09/724,676A

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: PatentIn version 3.2

SEQ ID NO 50957

LENGTH: 276
                                                       ; ORGANISM: Homo sapiens
US-09-724-676A-50957
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US-09-724-676A-50956
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US-09-724-676A-50956
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       Query Match
Best Local
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GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
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Best Local (
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LENGTH: 276
                                                                                        TYPE: PRT
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SOFTWARE: PatentIn version 3.2
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   Local Similarity
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APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: US/09/724,676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTMARE: Patentin version 3.2

EQ ID NO 50954

LENGTH: 328

TYPE: PRT

ORGANISM: Homo sapiens

US-09-724-676-50954
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US-09-724-676-50953
US-09-724-676-50953, Appli
GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patentin version 3.2
SEQ ID NO 50953
LENGTH: 328
TYPE: PRI
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902.626 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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	Score
100.0 97.4 96.9 96.9 96.9 96.9	Query Match Length DB
347 389 389 389 389	Length
2222244	BB
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Sequence 1, Appli Sequence 18, Appl Sequence 216, App Sequence 219, App Sequence 214, App Sequence 217, App	Description

Query Match 100.0%; Score 1929; DB 22; Length 347; Best Local Similarity 100.0%; Pred. No. 4.3e-141; Matches 347; Conservative 0; Mismatches 0; Indels 0;

0; Gaps

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ALIGNMENTS

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US-09-890-752A-1

Sequence 1, Application US/09890752A

GENERAL INFORMATION:

APPLICANT: Hildt, Eberhard

APPLICANT: Hofschneider, Peter

TITLE OF INVENTION: Particles for Gene Therapy

FILE REFERENCE: 107070-120 (VOS-013)

CURRENT APPLICATION NUMBER: US/09/890,752A

CURRENT FILING DATE: 2001-03-03

PRIOR FILING DATE: 2001-02-04

PRIOR APPLICATION NUMBER: DE 199 04 800.2

PRIOR APPLICATION NUMBER: DE 199 04 800.2

PRIOR FILING DATE: 1999-02-05

NUMBER OF SEQ ID NOS: 21

SOFTWARE: Patentin version 3.1

LENGTH: 347

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURES

FORMATION: Fusion protein comprising a LHBs and heterologous binding

US-09-890-752A-1
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APPLICANT: STUYVER Lieven
APPLICANT: VAN GEYT Caroline
APPLICANT: DE GENDT Sija
APPLICANT: DE GENDT Sija
APPLICANT: DE GENDT Sija
FILE REFERRNCE: 2551-52
CURRENT APPLICATION NEW HBV Sequences
FILE APPLICATION NUMBER: US/09/718,095
CURRENT FILING DATE: 2000-11-22
PRIOR APPLICATION NUMBER: E99870252.6
PRIOR APPLICATION NUMBER: US60/169,287
PRIOR APPLICATION NUMBER: US60/169,287
PRIOR FILING DATE: 1999-12-07
NUMBER OF SEQ ID NOS: 169
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 18
LENGTH: 389
TYPE: PAT
ORGANISM: Hepatitis B virus
US-09-718-095-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-718-095-18
US-09-718-095-18
; Sequence 18, Application US/09718095
; GENERAL INFORMATION:
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Matches 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VQWEVGLSPTVMLSVIMMMYWGPSLYSILSPELPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HSPTSCPPTCPGYRWMCLRRF11FLF1LLLCL1FLLVLLDYQGMLPVCPL1PGSSTTSTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HSPTSCPPTCPGYRWMCLRRPIIFLFILLLCLIFLLVLLDYQGMLFVCFLIPGSSTTSTG
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                                                                                                                                                                                                                                                                                                GAGAFGLGFTPPHGGLLGWSPQAQGIIQTLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 107
                                                                                                                                                                                                                                                                                                                   GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQFTPLSPPLRNTHPQ 65
                                                                                                                                                                                                                                                                                                                                                             97.4%;
cal Similarity 98.5%;
337; Conservation
                                                                                                                                                                                        TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSFTSNHSFTS 185
                                                                                                                                                                                                                                AMOMNISTTPHOTLODERVRGLYFPAGGSSSGTVNPVPTTASPISSIFSRIGDPALNMENI 167
                                                                                                 CPPTCPGYRMMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC 245
GLGPTVWLSVIWMWWWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                 TTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                     TTPAQGTSMYDSCCCTKDSDGNCTCIPIDSSWAFGKFLWEWASARFSWLSLLVDFVQWFV
                                                                                                                                                                     TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
                                                                                                                                                                                                                                                                                                                                                                    Score 1879; DB 21; Length 389; Pred. No. 3.7e-137; 2; Mismatches 3; Indels 0
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US-07-847-951A-216
US-07-847-951A-216
Sequence 216, Application US/07847951A
Sequence 216, Application US/07847951A
FOR THE PROJECT ON GENETICALLY ENGINEERED VACCINE TITLE OF INVENTION: STRAIN
FOR THE OF SEQUENCES: 462
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
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US-07-847-951A-216
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COURTES FORDERSEE: C/O William S. Frommer
STREET: 530 Fifth Avenue
CITY: New York
STATE: NY
COUNTRY: USA
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/847,951A
FILING DATE: 19920326
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REGISTRATION NUMBER: 25,506
REGISTRATION NUMBER: 454310-2400
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 040-3333
                                                                                                                                                                                                                                                                                                                                                                                           Query Match 96.9%;
Best Local Similarity 98.0%;
                                                                                                                                                                                                                                                                                                                                                                             Matches 335;
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TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 216:
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LENGTH: 389 amino acids
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STRANDEDNESS: single
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                                              MTTÄQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                                                                                 CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIFGSSTTSTGPCRTC
                                                                                                                                CPPTCPGYRMMCLRRFIIFLFILLLCLIFLLVLLDYQGMLFVCFLIFGSSTTSTGFCRTC
                                                                                                                                                                             TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
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RESULT 5
US-08-036-218-214
; Sequence 214, Application US/08036218
; GENERAL INFORMATION:
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Sequence 219, Application US/07847951A;
GENERAL INFORMATION;
APPLICANT: Paoletti, Enzo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/07/847,951A
PILING DATE: 19920326
CLASSIFICATION: 424
AITORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2400
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-333
INFORMATION FOR SEQ ID NO: 219:
SEQUENCE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO: 219:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 96.9%;
Best Local Similarity 98.0%;
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MEDIUM TYPE: Ploppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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CORRESPONDENCES: 462
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TYPE: AMINO ACID
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 530 Fit
CITY: New York
STATE: NY
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ADDRESSEE: c/o William S. Frommer
STREET: 530 Fifth Avenue
                                                                                                                       GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                       GLSPTVMLSVIMMMYMGPSLYSILSPFLPLLPIFFCLMVYI
                                                                                                                                                              MTTAQGTSMYPSCCCTKPSDCNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 347
                                                                                                                                                                                 TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFIWEWASARFSWLSLLVPFVQWFV 305
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 3; Mismatches

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APPLICATION NUMBER: US 07/666,056
FILING DATE: 11-JUN-1991
APPLICATION NUMBER: US 07/666,056
FILING DATE: 07-MAR-1991
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE,DOCKET NUMBER: 454310-22:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                      Matches 335;
                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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APPLICANT: Limbach, Keith J.
APPLICANT: Limbach, Keith J.
APPLICANT: Johnson, Gerard P.
APPLICANT: Johnson, Gerard P.
APPLICANT: Cox, William I.
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE STRAIN
NUMBER OF SEQUENCES: 221
CORRESPONDENCE ADDRESS:
ADDRESSEE: William S. Frommer c/o
ADDRESSEE: Curtis, Morris & Safford, P.C.
STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
288 MTTAQGTSMYPSCCCCTKPSDGNCTCIPIPSSWAPGKFLWEWASARPSWLSLLVPFVQWFV
                     246 TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARPSWLSLLVPFVQWFV 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                           168 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
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APPLICANT:
APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                                     66 AMQWNSTTFHOTLODFRVRGLYFFAGGSSSGTVNFVFTTVSFISSIFSRIGDFALNMENI 125
                                                                                                                                                                                                                                                                                                   48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
TOPOLOGY: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                            6 GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                        CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC
                                                                   CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC 287
                                                                                                                                                                TSGFLGPLLVLQAGEFLLTRILTIPOSLDSWWTSLNFLGGTTVCLGONSQSPTSNHSPTS 185
                                                                                                                                                                                                                AMQWNSTTPHQTLQDPRVRGLYFPAGGSSSGAVNPVPTTASPLSSIFSRIGDPALNMENI
                                                                                                                                                                                                                                                                                     GAGAFGLGFTPFHGGLLGWSFQAQGILQTLFANFPFASTNRQTGRQFTFLSFFLRNTHFQ 107
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: United State of America
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Norton, Elizabeth K.
Riviere, Michel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single
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Pred. No. 1.8e-136;
3; Mismatches 4; Indels
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GLSPTVWLSVIWMWYWGPSLYSILSPELPLLPIFFCLWVYI 389

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GENERAL INFORMA
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                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 217:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: not relevant
                                                                                                                                                                                                           Query Match
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APPLICANT:
TITLE OF IN
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APPLICANT:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
COMPUTER: PC COMPATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/713,967
FILING DATE: 11-JUN-1991
APPLICATION NUMBER: US 07/666,056
FILING DATE: 07-MAR-1991
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2291
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                  "108 AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGAVNPVPTTASPLSSIPSRIGDPALNMENI 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: New York
STATE: New York
COUNTRY: United State of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: William S. Frommer c/o
ADDRESSEE: Curtis, Morris & Safford, P.C.
STREET: 530 Fifth Avenue
126 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                      66
                                                                                                         48
                                                                                                                                                                       Local Similarity
les 335; Conserv
                                                                                                                                          o
                                                                                                       GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                        AMQMNSTTFHOTLODERVRGLYFBAGGSSSGTVNFVFTTVSFISSIFSRIGDFALMMENI 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10036
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de Taisne, Charles
Limbach, Keith J
Johnson, Gerard P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pincus, Steven B. Cox, William I.
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Norton, Elizabeth K.
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Taylor, Jill
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                                                                                                                                                                                                                                                                                                                                                                      (212) 840-0712
                                                                                                                                                                          Conservative
                                                                                                                                                                                           96.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US/08/036,218
                                                                                                                                                                              Score 1870; DB 4;
pred. No. 1.8e-136;
3; Mismatches 4;
                                                                                                                                                                                                               Length 389;
                                                                                                                                                                                  Indels
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                                                                                                           Matches 336; Conservative
                                                                                                                                                                                                                                                                          TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 116:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
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APPLICANT: Langlois, Alphonse J.

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION
TITLE OF INVENTION: OF MEMBRANE FUSION-ASSOCIATED EVENTS, INCLUDING HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      168
                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC COMPAtible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN PC-LOSS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/360,107
PILING DATE: 20-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                             NAME: COTUZZÍ, LBURB A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 78
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9990
TELEPHONE: (212) 790-9990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                  TOPOLOGY: unknown MOLECULE TYPE: protein
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STATE: New Yor
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                                                                                                                                                                                                                                                                                                                                              TELEFAX:
66 AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                          Local Similarity
                                                                                                                                                                                                                                    STRANDEDNESS:
                                     48
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                                                          GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                                                                                                                                                                                         amino acid
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Matthews, Thomas J.
Wild, Carl T.
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                                                                                                                                                                                                                                                                                                                                                (212) 869-9741/8864
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        York
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                                                                                                           96.8%; Score 1868; DB 7; Length 389; 98.2%; Pred. No. 2.6e-136; ative 2; Mismatches 4; Indels
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                                                                                                                     Gaps
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; MOLECULE TYPE:
US-08-470-896-106
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
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                                         Matches
                                                      Query Match 96.8%;
Best Local Similarity 98.2%;
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GENERAL INFORMATION:
APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
APPLICANT: Barney, Shawn O.
                                                                                                                                                                                                                                                                                                                                             NAME: COTUZZÍ, LAUTA A.
REGISTRATION NUMBER: 30,742
REPERENCE/DOCKET NUMBER: 7872-(
TELECOMMUNICATION INFORMATION;
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lambert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION
TITLE OF INVENTION: OF MEMBRANE FUSION-ASSOCIATED EVENTS, INCLUDING HIV
TITLE OF INVENTION: TRANSMISSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 1155 AVECTITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       348 GLSPTVWLSVIWMWYWGPSLYSILSPFLPLLPIFFCLWVYI 389
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                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               306 GLSPTVWLSVIWMMWYWGPSLYSILSPFLPILPIFFCLWVYI 347
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6 GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                       336; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARPSWLSLLVPFVQWFV 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TSGPLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 227
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                                                                                                                     protein
                      Score 1868; DB 8; Lenyum ...
Pred. No. 2.6e-136;
Pred. No. 4; Indels
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                                                                  Length 389;
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US-08-471-913-106
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                                                                                                                                                                                                                                                                                           ZIP: 10036-2711

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/471,913
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
                                                                                                                       TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
           STRANDEDNESS:
TOPOLOGY: unk
MOLECULE TYPE:
                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: COCUZZI, LAURA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKST NUMBER: 7872-030
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9990
                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Barney, Shawn O.
APPLICANT: Lambert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF MEMBRANE
TITLE OF INVENTION: FUSION-ASSOCIATED EVENTS, INCLUDING EPSTEIN-BARR VIRUS
TITLE OF INVENTION: TRANSMISSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          228 CPPTCPGYRWMCLRRFIIFLFILLICLIFLLVILDYQGMLPVCPLIPGSSTTSTGPCRTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
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                                 unknown
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Matthews, Thomas J.
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US-08-475-668-106
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Matches 336; Conservative
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                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: LEM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/475,668

FILING DATE: 07-JUN-1995

CLASSIFICATION: 530

ATTORNEY/AGENT INFORMATION:

NAME: COTUZZI, LAURA A.

REGISTRATION NUMBER: 30,742

REFERENCE/DOCKET NUMBER: 30,742
                                                                                                                                                                                        TELEFAX: (212) 869-9741/886:
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
APPLICANT: Barney, Shawn O.
APPLICANT: Lambert, Dennis M.
APPLICANT: Lembert, Dennis M.
APPLICANT: Lembert, Stephen R.
APPLICANT: Lambort, Stephen R.
APPLICANT: Lambort, Alphonse J.
APPLICANT: Lambort, Stephen R.
APPLICANT: Lambort, Stephen R.
APPLICANT: Lambort, Stephen R.
APPLICANT: Lambort, Stephen R.
APPLICANT: Lambort, Alphonse J.
APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
APPLICANT: Wild, Carl T.
APPLICANT: Lambort, Dennis J.
APPLICANT: Lambort, Den
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                                                                                                                   SEQUENCE CHARACTERISTICS:
- LENGTH: 389 amino acids
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                                    STRANDEDNESS:
TOPOLOGY: unknown
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Matthews, Thomas J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                   (212) 869-9741/8864
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2; Mismatches 4;
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Sequence 106, Application US/08484223;
GENERAL INFORMATION:
APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Mild, Carl T.
APPLICANT: Barney, Shawn O.
APPLICANT: Lambert, Dennis M.
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US-08-475-668-106
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Best Local Similarity 98.7
Matches 336; Conservative
                                                                                                                                                                                                    CCUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/484,223
FILING DATE: 07-JUN-1995
TELEPHONE: (212) 790-9090
TELEPAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
                                                                                     CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, Laura A.
REGISTRATION UNMER: 30,742
REFERENCE/DOCKET NUMBER: 7872
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF
TITLE OF INVENTION: MEMBRANE FUSION-ASSOCIATED EVENTS, INCLUDING
TITLE OF INVENTION: TRANSMISSION
NUMBER OF SEQUENCES: 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               168 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSFTSNHSPTS
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                                                                                                                                                                                                                                                                                                                                                                                           STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Pennie &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLQQNSQSPTSNHSPTS 185
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                                                                                                                                                                                                                                                                                                                                                                                             New York
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1155 Avenue of the Americas
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98.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 1868, DB 8;
Pred. No. 2.6e-136;
                                                                                                                    7872-029
                                                                                                                                                                                                                                                                  Version
                                                                                                                                                                                                                                                                      #1.30
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SEQUENCE 106, Application US/08484223A

APPLICANT: Bolognesi, Dani P.

APPLICANT: Matthews, Thomas J.

APPLICANT: Wild, Carl T.

APPLICANT: Barney, Shawn O.

APPLICANT: Barney, Shawn O.

APPLICANT: Lambert, Dennis M.

APPLICANT: Lambert, Dennis M.

APPLICANT: Langlois, Alphonse J.

APPLICANT: Langlois, Alphonse J.

TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF

TITLE OF INVENTION: MEMBRANE FUSION-ASSOCIATED EVENTS, INCLUDING HIV

NUMBERS OF STATEMENTS.
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COMPUTER READBLE FOORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNMER: US/08/484,223A
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-029
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 96.8%;
Best Local Similarity 98.2%;
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                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2. CORRESPONDENCE ADDRESS:
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ZIP: 10036-2711
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TOPOLOGY: un
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RESULT 13
US-08-485-546-106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 106, Application:
ZIP: 10036-2711

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Ve:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,546
PILING DATE: 07-JUN-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAURA A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 336; Conserv
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acid
                                                                                                                                                                                                                                                                                                   APPLICANT: Lambert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: METHODS FOR INHIBITION OF MEMBRANE
TITLE OF INVENTION: PUSION-ASSOCIATED EVENTS, INCLUDING
TITLE OF INVENTION: TRANSMISSION
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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                                                                                                                                                                                                              CITY: New York
STATE: New York
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                                                                                                                                                                                                                                                               ADDRESSEE:
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TYPE: amino acid
STRANDEDNESS:
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                                                                                                                                                                                                                                                1155 Avenue
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                                                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                          Barney, Shawn O.
Lambert, Dennis M.
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Pred. No. 2.6e-136;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                             Version #1.30
                                                                                                                                                                                                                                                                                                                        INCLUDING EPSTEIN-BARR VIRUS
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; MOLECULE TYPE: protein US-08-485-546-106
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Sequence 106, Application US/08485546A
GENERAL INFORMATION:
APPLICANT: Bolognesi, Dani P.
APPLICANT: Bolognesi, Dani P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                문
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Best Local Similarity 98.2%;
Matches 336; Conservative
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
COUNTRY: USA
ZIP: 1036-2711
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PACENTIA PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,546A
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REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
APPLICANT: Barnev ch-
                                                                                                                                                                                                                                                                   APPLICANT: Lambert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
APPLICANT: LANGLOIS, Alphonse J.
TITLE OF INVENTION: METHODS FOR INHIBITION OF MEMBRANE
TITLE OF INVENTION: FUSION_ASSOCIATED EVENTS, INCLUDING EPSTEIN-BARR VIRUS
TITLE OF INVENTION: TRANSMISSION
NUMBER OF SEQUENCES: 214
                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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CITY: New York
STATE: New York
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TOPOLOGY: un
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 389 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6 GAGAPGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                                                                                                                                                                   ADDRESSEE: Pennie & Edmonds LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAGAFGLGFTPPHGGLLGWSPQAQGILQTLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CPPTCPGYRWMCLRRF11FLF1LLLCL1FLLVLLDYQGMLPVCPL1PGSSTTSTGPCRTC
                                                                                                                                                                           New York
                                                                                                                                                                                                               1155 Avenue of the Americas
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Pred. No. 2.6e-136;
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; MOLECULE TYPE: protein US-08-485-546A-106
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US-08-485-551-106
                                                                                                                                                                       Seguence 106, Application US/0848551

GENERAL INFORMATION:
APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Mild, Carl T.
APPLICANT: Barney, Shawn O.
APPLICANT: Liambert, Dennis M.
APPLICANT: Liambert, Dennis M.
APPLICANT: Langlois, Alphonse J.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: METHODS FOR INHIBITION OF MEMBRANE
TITLE OF INVENTION: TRANSMISSION
NUMBER OF SEQUENCES: 209
NUMBER OF SEQUENCES: 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 96.8%;
Best Local Similarity 98.2%;
Matches 336; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/886
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: COTUZZI, LAURA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 78
TELECONMUNICATION INFORMATION:
TELEPHONE: (212) 790-9990
TELEPHONE: (212) 790-9990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZÍ, Laura A.
COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       348 GLSPTVWLSVIWMYWGPSLYSILSPFLPLLPIFFCLWVYI 389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           306 GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   246 TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     168 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSFTSNHSPTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            288 MTTAOGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 66 AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSBISSIFSRIGDPALMMENI 125
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                                                                                                       STREET: 1155 A
                                                                                          STATE: New York
                                                                                                                                              ADDRESSEE:
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1155 Avenue of the Americas
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Pred. No. 2.6e-136;
2; Mismatches 4;
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ATTORNEY/AGENT INFORMATION:

NAME: COTUZZI, LBUIRA A.

REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-023
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
STANDENMESS:
TOPOLOGY: unknown
MOLECULE TYPE: protein
US-08-485-551-106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,551
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
CLASSIFICATION: 514
246 TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
                                                                                                                                         186 CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC 245
                                                                                                                                                                                                              126 TSGELGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSFTSNHSPTS 185
                                                                                                                                                                                                                                                                                    66 AMOMNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                                                                                                                                                                                                                                                        6 GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
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Search completed: February 3, 2003, 09:34:07 Job time: 250.857 secs

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Result
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Maximum DB :
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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1721
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seq length: 2000000000
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1: /cgn2 6/ptodata/2/pubpaa/USO8 NEW PUB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/USO8 NEW PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/USO6 NEW PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/USO6_NEW PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/USO6_PUBCOMB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*

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9: /cgn2_6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*

9: /cgn2_6/ptodata/2/pubpaa/USO9_PUBCOMB.pep:*

10: /cgn2_6/ptodata/2/pubpaa/USO9_PUBCOMB.pep:*

10: /cgn2_6/ptodata/2/pubpaa/USO9_PUBCOMB.pep:*

11: /cgn2_6/ptodata/2/pubpaa/USO9_PUBCOMB.pep:*

12: /cgn2_6/ptodata/2/pubpaa/USO9_PUBCOMB.pep:*

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Match
             122226 seqs, 20178551 residues
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Copyright (c) 1993 - 2003 Compugen Ltd.
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9 US-09-247-890-12

9 US-09-247-890-10

10 US-09-247-890-10

10 US-09-221-877-8

10 US-09-821-877-8

10 US-09-821-862-14

10 US-09-812-862-4

10 US-09-812-862-4

10 US-09-812-862-4

10 US-09-812-862-2

10 US-09-812-866-54

10 US-09-818-066-54

10 US-09-818-066-54

10 US-09-818-066-46
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Sequence 2, Appli
Sequence 3, Appli
Sequence 10, Appl
Sequence 10, Appl
Sequence 10, Appli
Sequence 10, Appli
Sequence 8, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 16, Appli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09821877

Patent No. US20020177124A1

GENERAL INFORMATION:
APPLICANT: Abbott Laboratories
APPLICANT: Coleman, Paul F.
APPLICANT: Coleman, Paul F.
APPLICANT: Mushahwar, Isa K.
ITITLE OF INVENTION: Hepatitis B Virus Surface Antigen Mut
ITITLE OF INVENTION: And Methods Of Detection Thereof
FILE REFERENCE: 6794.US.01

CURRENT APPLICATION NUMBER: US/09/821,877

CURRENT APPLICATION ON THE COUNTY OF THE SEQ ID NOS: 8

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1
US-09-821-877-2
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TYPE: PRT
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ALIGNMENTS

Mutant

	246 TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARPSWLSLLVPPVQWFV 305	Оу 246	_
	CPPTCPGYRMMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRAC 287	Db 228	н
	186 CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLFVCPLIPGSSTTSTGPCRTC 245	Оу 186	_
	TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 227	Db 168	
	126 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185	Ωу 120	_
	AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPARNMENI 167	Db 108	_
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		Db 48	_
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0,	è	Matche	
	Query Match 96.7%; Score 1866; DB 9; Length 389; Best Local Similarity 98.2%; Pred. No. 1.9e-132;	Query Match Best Local	
	ORGANISM: Hepatitis B Virus 99-821-877-2	GRGANISM: H US-09-821-877-2	

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FILING DATE: 19-JAW-1998
ATTORNEY/AGENT INFORMATION:
NAME: MASS. Clifford J.
REGISTRATION NUMBER: 30,086
REFERENCE/DOCKET NUMBER: U-013109-7
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 708-1800
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 400 amino acids
TYPE: amino acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 3, Application US/10209264 Publication No. US20030003111A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 308; Conserv
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                             186
                                                                                                                                   119
                                                                                                126
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Ladds & Parry
STREET: 26 West 61 Street
CITY: New York
STATE: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTMARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/209,264
PILING DATE: 31-Jul-2002
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Oon, Chong Jin
Lim, Gek Keow
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: A MUTANT HUMAN HEPATITIS B VIRAL STRAIN
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                                                                                                                                                  AMOMNSTTEHOTLODDRVRGLYPPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                                                                                                 GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                              TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                TSGPLGPLLVLQAGPPSLTRILTIPQSLDSWWTSLNPLGGAPTCPGQNSQSPTSNHSPTS
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                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
                                                                                                                                                                                                                                                                89.2%; Score 1721; DB 9; ilarity 90.1%; pred. No. 1.3e-121; Conservative 10; Mismatches 24;
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                                                                       238
      298
                                    245
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APPLICANT: MANDAMANANA JUNA
APPLICANT: Bass, Steven H.
APPLICANT: Whalen, Robert Gerald
APPLICANT: Whalen, Robert Gerald
APPLICANT: Howard, Russell
APPLICANT: Howard, Russell
APPLICANT: Maxygen, Inc.
APPLICANT: Maxygen, Inc.
ITILE OF INVENTION: Antigen Library Immunization
FILE REFERENCE: 018097-028710US
FULE REFERENCE: 018097-028710US
CURRENT APPLICATION NUMBER: US 60/074,890
CURRENT FILING DATE: 1999-02-11
EARLIER APPLICATION NUMBER: US 60/074,294
EARLIER APPLICATION NUMBER: US 60/105,509
EARLIER APPLICATION NUMBER: US 60/105,509
EARLIER PILING DATE: 1998-10-23
NUMBER OF SEQ ID NO 12
SEQ ID NO 12
LENGTH: 281
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US-09-247-890-12
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US-09-247-890-12
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                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 276; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
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187
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                                                                                 TPAQGTSMYPSCCCTKESDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFVG
                                                                                                                                                 PPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTCT 246
                                                                                                                                                                                          SGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTSC 186
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98.2%;
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                                                                                                                                                                                                                                                                                                                         Score 1534; DB 9;
pred. No. 8.5e-108;
1; Mismatches 4;
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                                                                         240
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Sequence 10, Application US/09247890
Publication No. US20020198162A1
GENERAL INFORMATION:
APPLICANT: Punnonen, Juha
APPLICANT: Bass, Steven H.
APPLICANT: Whalen, Robert Gerald
APPLICANT: Howard, Russell
APPLICANT: Stemmer, Willem P.C.
APPLICANT: Maxygen, Inc.

US-09-247-890-10

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APPLICANT: Matti Sallberg

APPLICANT: Catharina Hultgren

ITITLE OF INVENTION: VACCINES CONTAINING RIBAVIRIN AND

ITITLE OF INVENTION: WATHOUS OF USE THEREOF

ITITLE OF INVENTION: WETHOUS OF USE THEREOF

CURRENT APPLICATION NUMBER: US/10/104,966

CURRENT FILING DATE: 2002-03-22

PRIOR APPLICATION NUMBER: 09/705,547

PRIOR APPLICATION NUMBER: 09/705,547

PRIOR APPLICATION NUMBER: 09/229,175

PRIOR APPLICATION NUMBER: 60/229,175

PRIOR APPLICATION NUMBER: 60/229,175

PRIOR FILING DATE: 2000-08-29

NUMBER OF SEQ ID NOS: 15

SOFTMARE: FASTSEQ for Windows Version 4.0

LENGTH: 2266
                                      5
                                                                                                                                             ; OTHER INFORMATION: Hepatitis B virus S antigen (HBsAg) sequence
US-10-104-966-10
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                                                           Query Match 65.3
Best Local Similarity 99.1
Matches 224; Conservative
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; ORGANISM: Hepatitis B virus
US-09-247-890-10
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SEQ ID NO 10
LENGTH: 281
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                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Antigen Library Immunization FILE REPERENCE: 018097-028710US
122 MENITSGELGELLVLQAGEFLLTRILTIPQSLDSWWTSLNELGGTTVCLGQNSQSPTSNH 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           241
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                                                                                  65.3%;
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91.5%;
                                                         Score 1259; DB 9;
Pred. No. 2.4e-87;
0; Mismatches 2;
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Pred. No. 1.5e-100;
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                                                                                             Length 226;
                                                               Indels
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Best Local S
Matches 224
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CURRENT FILING DATE: 2001-08-15
PRIOR APPLICATION NUMBER: 09/705,547
PRIOR PILING DATE: 2000-11-03
PRIOR APPLICATION NUMBER: 60/229,175
PRIOR FILING DATE: 2000-08-29
PRIOR APPLICATION NUMBER: 60/225,767
PRIOR APPLICATION NUMBER: 60/225,767
PRIOR FILING DATE: 2000-08-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Matti Sallberg
APPLICANT: Catharina Hultgren
TITLE OF INVENTION: VACCINES CONTAINING RIBAVIRIN
TITLE OF INVENTION: METHODS OF USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: TRIPEP.23AUS2
                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Hepatitis B virus S antigen (HBsAg) sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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                                                                  181
                                                                             302 QWEVGLSPTVWLSVIWWWWYWGPSLYSILSPELPLLPIFECLWVYI 347
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                                                                                                                                                                                                                                                      1 MENITSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNH
                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                        QWFVGLSPTVWLSVIWMWYWGPSLYSILSPFLPLLPIFFCLWVYI 226
                                                                                                                       CRTCMTTĂQGTŚMYPSCCCCTKPSDGNCTCIPIPSSWAPGKFLWEWASARFŚWLSLLVPFV
                                                                                                                                             CRTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFV 301
                                                                                                                                                                                       SPTSCPPTCPGYRWMCLRRPTIFLFILLLCLIPLLVLLDYQGMLPVCPLIPGSSTTSTGP
                                                                                                                                                                                                         SPTSCPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGP
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                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                         65.3%; Score 1259; DB 10; 99.1%; Pred. No. 2.4e-87; tive 0; Mismatches 2;
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Sequence 8, Application US/09821877 Patent No. US20020177124A1 GENERAL INFORMATION:

APPLICANT: Abbott Laboratories
APPLICANT: Coleman, Paul F.
APPLICANT: Mushahwar, Isa K.
TITLE OF INVENTION: Hepatitis!

W

Virus Surface Antigen Mutant

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SEQ ID NO 8
LENGTH: 229
TYPE: PRT
ORGANISM: Hepatitis B Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
NAME/KEY: VARIANT
LOCATION: (126)...(126)
OTHER INFORMATION: Xaa = A ON
NAME/KEY: VARIANT
LOCATION: (202)...(202)
OTHER INFORMATION: Xaa = L ON
NAME/KEY: VARIANT
LOCATION: (210)...(210)
OTHER INFORMATION: Xaa = T ON
OTHER INFORMATION: Xaa = T ON
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CURRENT APPLICATION MUMBER: US/09/821,877
CURRENT FILING DATE: 2000-03-03
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Application US/09812862
Patent No. US20020035081A1
GENERAL INFORMATION: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        119 ALNMENITSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPT 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
hes 225; Conserv
                                                         COUNTRY: USA
ZIF: 02110-2804
ZIF: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Plom PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Ver
SOPTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/812,862
FILING DATE: 20-MAY-2001
CLASSIFICATION SYMBER: US/09/812,862
PRIOR APPLICATION NUMBER: US/09/812,862
                                                                                                                                                                                                                                                                                                                                                                      Scaglioni, Pier Paolo
Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PPVQWEVGLSPTVWLSVIWMMYWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNHSPTSCPPTCPGYRWMCLRRFIIPLPILLLCLIPLLVLLDYQGMLPVCPLIPGSSTTS 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNHSPTSCPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTS 238
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGPCRTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLV 298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PEVOMEVGLSPTVMLSVIMMXYMGPSLYXILSPFLPLLPIFFCLMVYI 229
                                                                                                                                                                                                                                                                                                                               CITY: Boston
STATE: MA
APPLICATION NUMBER: 08/667,073 FILING DATE: <Unknown> APPLICATION NUMBER: 08/492,489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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Pred. No. 2.5e-87;
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SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-09-812-862-14
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Best Local Similarity
Matches 223; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6, Application US/09812862
Patent No. US20020035081A1
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 14:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 20-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 0078
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5970
                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PACENTIN Release #1.0, Ver

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/812,862

FILING DATE: 20-Max-2001

CLASSIFICATION: <Unknown>

PRIOR APPLICATION NUMBER: 08/667,073

PILING DATE: <Unknown>

APPLICATION NUMBER: 08/492,489

FILING DATE: 20-JUN-1995

FILING DATE: 20-JUN-1995

FILING DATE: 20-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 SPTSCPPTCPGYRWMCLRRP11F1F1LLLCL1F1LVLLDYQGMLPVCPL1PGSSTTSTGP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MENITSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Wands, Jack R
                                       ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPAX: 617/542-8906
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                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
                    REGISTRATION NUMBER: 30,162
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pred. No. 8.1e-87;
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                                                                                                                                                                                                                                                                                          Version #1.30
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Query Match

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US-09-812-862-14

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RESULT 10
US-09-812-862-4
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' SEQUENCE DESCRIPTION: SEQ ID NO: 6:
US-09-812-862-6
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/09812862 Patent No. US20020035081A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 63.8%;
Best Local Similarity 69.7%;
Matches 237; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 200154
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                             ZIP: 02110-2804

ZIP: 02110-2804

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/812,862

FILING DATE: 20-Mar-2001

CLASSIFICATION DATA:

ARBITCATION DATA:

ARBITCATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             128 GPLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSBTSCP 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            121 SPGVWIRTPP----AYRPPNAPILSTL----PETTVVRRRGRSPRRRTPSPRRRRSQS- 170
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                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                        Scaglioni, Pier Paolo
Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLMVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARPSWLSLLVPPVQWFVGL 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAPGKPLWEWASARFSWLSLLVPFVQWFVGL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PTCPGYRMMCLRRFIIFLFILLCLIFLLVLLDYQGMLFVCFLIFGSSTT8TGFCRTCTT 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIFGSSTTSTGPCRTCMT 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENITS 127
              APPLICATION NUMBER: 08/667,073 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                       CITY: Boston
STATE: MA
APPLICATION NUMBER: 08/492,489
                                                                                                                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
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Pred. No. 5.6e-85;
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GENERAL INFORMATION:

APPLICANT: Bass, Steven H.

APPLICANT: Bass, Steven H.

APPLICANT: Howard, Russell

APPLICANT: MAXYSEN, Inc.

TITLE OF INVENTION: Antigen Library Immunization

FILE REFERENCE: 018097-02871005

FURBERT APPLICATION UMMBER: US/09/247,890

CURRENT FILING DATE: 1999-02-10

EARLIER APPLICATION NUMBER: US 60/074,294

EARLIER APPLICATION UMMBER: US 60/105,509

INMBER OF SEQ ID NOS: 25

SOFTWARE: PATENTING DATE: 1998-10-23

NUMBER OF SEQ ID NOS: 25

SOFTWARE: PATENTING DATE: 1998-10-23

LENGTH: 282

TYPE: DPT
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                                                                                                                                                                      Query Match 43.3%; Score 834.5; DB 9; Best Local Similarity 56.8%; Pred. No. 1.7e-55; Matches 159; Conservative 29; Mismatches 85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16, Application US/09247890 Publication No. US20020198162A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity Matches 174; Conserv
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                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Woodchuck hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 200154
INFORMATION FOR SEQ ID NO: 4:
                      128 GPLGPLLVLQAGEFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSDTSNHSDTSCP 187
             5
                                                                                                                         73 TFH-----QTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALMWENITS 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       176 ONSOSPTSNHSPTSCPPTCPGYRWMCLRRPTIFLFILLLCLIFLLVLLDYQGWLPVCPLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 292 SWLSLLVPFVQWFVGLSPTVWLSVIWMMYWGPSLYSILSPELPLLDIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  236 PGSSTTSTGPCRTCMTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               232 PGSSTTSTGPCRTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKPLWEWASARF 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  172 ONSQSETSNHSETSCPETCEGYRWMCLERFEIIFLEILLICLIFLLVLLDYQGMLEVCPLI 231
                                                                                                   u
SLIGLLAGLÓVVYFLMTKILTIAQNI.DWWWTSLSFPGGIPECTGONSOFOTCKHLFTSCP 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 351 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                   TFHLQGFVDGLRDLTTTERQHNAYGDPFTTLSPAVPTVSTILSPPSTTGDPALSPEMSPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 20-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Clark, Paul T. REGISTRATION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617/542-8906
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98.9%;
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Pred. No. 1.5e-67;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 00786/282001
                                                                                                                                                                                                                 Length 282;
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                                                                                                                                                                      Gaps
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US-09-812-862-8
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Patent No. US20020035081A1
GENERAL INFORMATION:
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Best Local (
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INFORMATION FOR SEQ ID NO: 8:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC COMPACTIBLE
COMPUTER: IBM PC COMPACTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.30
SOFTMARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/812,862
PILING DATE: 20-Mar-2001
CLASSIPICATION UMBER: US/09/812,862
PILING DATE: UNKNOWN>
PRIOR APPLICATION NUMBER: 08/667,073
APPLICATION NUMBER: 08/667,073
APPLICATION NUMBER: 08/667,073
APPLICATION UMBER: 08/667,073
APPLICATION UMBER: 08/667,073
APPLICATION UMBER: 08/492,489
                      . 294 LSLLVPEVQMEVGLSPTVMLSVIMMMYMYWGPSLYSILSPELPLLPIFECLMVYI 347
                                                                                    176
                                                                                                             234 SSTTSTGPCRTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSW 293
                                                                                                                                                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scaglioni, Pier Paolo
Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SPTVWLSVIWMWWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFVGL 307
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/282001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5970
                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                    SSTTSTGPCRTCMTTAQGTSMYPSCCCTKPSDGNCTC1P1PSSWAPGKFLWEWASARESW 235
STATE: MA
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                                                                                                                                                                                                                                                                                                                                                 LENGTH: 289 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 617/542-8906
                                                                                                                                                                           Conservative
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                                                                                                                                                                        Score 652; DB 10;
pred. No. 7.5e-42;
0; Mismatches 2;
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RESULT 13
US-09-812-862-2
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Patent No. US20020035081A1
GENERAL INFORMATION:
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RESULT 14
US-09-818-066-55
; Sequence 55, Application US/09818066
; Patent No. US20020032307A1
; GENERAL INFORMATION:
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                  172 QNSQSPTSNHSPTSCPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLI 231
                                                                                                                                                                                                                                                                               173
                                                                                                                                                                                                         233
                                                                                                                                                             292 SMISLIVPEVOMEVGLSPTVMISVIMMYMGPSLYSILSPELELLEIFECLMVYI 347
                                                                                                                                291
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ADDRESSEE: Fish & Richardson
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Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/667,073
FILING DATE: <Unknown>
APPLICATION NUMBER: 08/492,489
FILING DATE: 20-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/282001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEPAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                         PGSSTTSTGPCRTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLMEWASARF 291
                                                                                                                                                                                                                                                                             PTTETTVN--CROCTISACOMYTPPYCCCLKPTAGNCTCWPIPSSWALGNYLWEWALARL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/812,862 FILING DATE: 20-Mar-2001 CLASSIFICATION: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: .225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 346 amino acids
                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                      33.6%; Score 649; DB 10; 63.1%; Pred. No. 1.5e-41; tive 21; Mismatches 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                Length 346;
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APPLICANT: Shuping Tong et al

TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR NUMBER OF SEQUENCES: 75

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RESULT 15
US-09-818-066-54
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                                                                                                                                                                                                                      Sequence 54, Application US/09818066
Patent No. US20020032307A1
GENERAL INFORMATION:
APPLICANT: Shuping Tong et al.
TITLE OF INVENTION: HEPADNAVIRUS RECEPTOR NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS;
ADDRESSES: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-818-066-55
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 55: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 119 AMQMNSTTFHOTLODÞRVRGLÍFÞAGGSSSGTVNÞVÞTTASÞÍSSÍÞSRÍGÐÞÁLN 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           66 AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALN 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       59
COMPUTER READABLE FORM:

MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION NUMBER: US/09/818,066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 GAGAPGIGFTPPHGGILGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 55:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/818,066
PILING DATE: 27-Mar-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/683,262
PILING DATE: 18-UUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
REFERENCE/DOCKET NUMBER: 00786/287002
TELECOMMUNICATION INFORMATION:
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ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                         STATE: Massachusetts
                                                                                                                                                                                             CITY: Boston
                                                                                                                                                            COUNTRY: U.S.A.
                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 174 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
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STATE: Massachusetts
COUNTRY: U.S.A.
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                                                                                                                                                                                                               225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31.8%; Score 613; DB 10;
96.6%; Pred. No. 3.5e-39;
tive 2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 174;
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                                                                                                                                       Best Local Similarity
Matches 112; Conservat
                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                     TELEX: 200154
INFORMATION FOR SEQ ID NO: 54:
119 AMOMNSTTEHQTLODPRVRGLYLPAGGSSGGTVNPVPTTVSPISSIFSRTIDPALN 174
                   66 AMQWNSTTFHQTLQDPRVRGLYFDAGGSSGGTVNPVPTTVSPISSIFSRIGDPALN 121
                                                                            59
                                                                                                           0
                                                                                                                                                                                                              LENGTH: 174 amino acids
TYPB: amino acid
TYPB: amino acid
TOPOLOGY: linear
MOLECULE TYPB: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 54:
                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/683,262
FILING DATE: 18-UTU-1996
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 31,819
                                                           GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 27-Mar-2001
CLASSIFICATION: <Unknown
                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617) 542-8906
                                                                                                                                       Conservative
                                                                                                                                                     31.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <Unknown>
                                                                                                                                     <u>,</u>
                                                                                                                                Score 604; DB 10;
Pred. No. 1.7e~38;
1; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                     00786/287002
                                                                                                                                                               Length 174;
                                                                                                                                  Indels
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Search completed: February 3, 2003, 09:35:27 Job time: 13.6951 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.

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Match
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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Gapop 10.0 , Gapext 0.5
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1929
                  262574 seqs, 29422922 residues
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                                                                                                                                                                                                                                                                                                                                                                                         Length
                  GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd
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482.536 Million cell updates/s
             US-08-105-483-216
US-08-105-483-219
US-08-709-209-219
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US-08-458-101-216
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US-08-458-107A-116
US-08-485-23B-106
US-08-475-668A-106
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US-08-471-913A-106
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US-08-471-913A-106
US-08-471-913A-106
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US-08-793-29-214
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US-08-793-29-214
US-08-793-29-214
US-08-793-29-214
US-08-793-29-29-1
US-08-793-29-29-1
US-08-793-29-29-3
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      Sequence 216, App Sequence 219, App Sequence 211, App Sequence 211, App Sequence 216, App Sequence 216, App Sequence 106, App Sequence 216, App Sequence 214, App Sequence 214, App Sequence 214, App Sequence 214, App Sequence 11, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 3, Appli Sequence 6, Appli
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GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ
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Result No.

Pred. No. score grea and is der

27653211654 27652211654 27652211654 27652211654

Minimum Maximum

88

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Scoring table:

Title: Perfect score:

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protein -8

Database

Ouery Match 96.9%; Score 1870; DB 1; Length Best Local Similarity 98.0%; Pred. No. 3.4e-157; Matches 35; Conservative 3; Mismatches 4; Indel	Sequence 216, Application US/08105483 Sequence 216, Application US/08105483 PATERI NO. 5494807 GENERAL IMPORMATION: APPLICANT: PAOLETT! RAIN APPLICANT: PAOLETT! STRAIN INTHE OF INVERVION: GENETICALLY ENGINEERED VACCINE TITLE OF INVERVION: GENETICALLY ENGINEERED VACCINE TITLE OF INVERVION: GENETICALLY ENGINEERED VACCINE TITLE OF INVERVION: 462 CORRESPONDENCE ADDRESS: ADDRESSEE: C/O William S. Frommer STREET: 530 Pitch Avenue CITY: New York STATE: NY COUNTRY: USA ZIP: 10036 COMPUTER READABLE FORM: MEDIUM TYPE: Ploppy disk COMPUTER: IBM PC compactible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION NATA: APPLICATION NUMBER: US/08/105,483 FILING DATE: 12-AUG-1993 CLASSIFICATION: 42 APPLICATION NUMBER: US/07/847,951 FILING DATE: 06-MAR-1992 APTORNEY/AGENT INFORMATION: APPLICATION NUMBER: 25,506 REFERENCES/DOCKET NUMBER: 454310-2400 TELEPHONE: (212) 840-2333 INFORMATION FOR SEQ ID NO: 216: SEQUENCE CHARACTERISTICS: LENGTH: 389 amino acids STRANDEDNESS: single TOPOLOGY: linear TOPOLOGY: linear	ALIGNMENTS	29 1217 63.1 226 6 5196194-21 29 1217 63.1 226 6 5436139-5 30 1212 62.8 225 6 5436139-4 31 1206 62.5 226 1 US-08-378-011A-3 32 1197 62.1 226 6 5198348-1 33 1160 60.1 228 1 US-08-447-591-2 34 1160 60.1 228 1 US-08-450-943-2 35 1160 60.1 228 1 US-08-450-942-2 36 1160 60.1 228 1 US-08-450-942-2 37 1160 60.1 228 2 US-08-450-942-2 38 1142 59.2 228 1 US-08-450-942-3 39 1142 59.2 228 1 US-08-450-943-3 40 1142 59.2 228 1 US-08-450-943-3 41 1142 59.2 228 1 US-08-059-031-3 42 1142 59.2 228 2 US-08-450-943-3 43 1114 57.8 228 1 US-08-450-943-3 44 1114 57.8 228 1 US-08-450-943-3 45 1114 57.8 228 1 US-08-450-943-4 46 1114 57.8 228 1 US-08-450-943-4 47 114 57.8 228 1 US-08-450-943-4
389; 8 0;			Patent Mo Patent No. Patent No. Sequence Patent No. Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence
Gaps 0;			. 5196194 . 5436139 . 5436139 . 3, Appli . 5, Appli . 5, Appli . Appli

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RESULT 2
US-08-105-483-219
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                                                                                                                                       APPLICATION NUMBER: US/08/105,483
FILING DATE: 12-AUG-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/847,951
FILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2400
FELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEPHONE: (212) 840-3333
TELEPHONE: (212) 840-319:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
TYPE: amino acids
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 219, Application US/08105483 Patent No. 5494807
                                                    Best Local Similarity Matches 335; Conserv
                                                                                           Query Match
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APPLICANT: Paoleti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
TOTAL TATALY NATION. DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: GE
TITLE OF INVENTION: ST
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 550
STREET: 550
CITY: New York
CTATE: NY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Curtis, Morris & Safford ADDRESSEE: c/o William S. Frommer STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GLSPTVWLSVIWMMYWGPSLYSILSPFLPLLPIFFCLMVYI 389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAGAFGLGFTPPHGGLLGWSPQAQGILQTLPANPPPASTNRQTGRQPTPLSPPLRNTHPQ 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Paoletti, Enzo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENETICALLY ENGINEERED VACCINE STRAIN
                                                                          96.9%;
98.0%;
                                                         ω
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                                                         Score 1870; DB 1;
Pred. No. 3.4e-157;
3; Mismatches 4;
                                                                                           Length 389;
                                                           Indels
                                                           0;
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                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/709,209
FILING DATE: 21-AUG-1996
FILING DATE: 21-AUG-1996
FILING DATE: 12-AUG-1996
APPLICATION NUMBER: US/07/847,951
FILING DATE: 12-AUG-1993
APPLICATION NUMBER: US/07/847,951
FILING DATE: 12-AUG-1992
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REGISTRATION NUMBER: 454310-2400
TELEPHONE: (212) 840-3333
TELEPHONE: (212) 840-3333
TELEPAX: (212) 840-3333
TELEPAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 216:
SEQUENCE CHARACTERSTICS:
LENGTH: 389 amino acide
CTEANUPPINES: atingle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-709-209-216
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Patent No. 5762938
GENERAL INFORMATION:
Query Match 96.9%; Score 1870; DB 1; Best Local Similarity 98.0%; Pred. No. 3.4e-157; Matches 335; Conservative 3; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: PAOLETTI, ENZO
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           186
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ADDRESSEE: C/O William S. Frommer
STREET: 530 Fifth Avenue
CITY: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                            STRANDEDNESS:
TOPOLOGY: li
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                                                                                                          linear
                                                                                                                             single
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                                                   Length 389;
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Indels

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Gape

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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US 08/709,209
FILING DATE: 21-AUG-1996
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: US 08/105,483
FILING DATE: 12-AUG-1993
APPLICATION NUMBER: US 07/847,951
FILING DATE: 12-AUG-1993
APPLICATION NUMBER: US 07/847,951
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 44,310-2400
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-313
TELEPAX: (212) 840-313
INFORMATION FOR SEQ ID NO: 219:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 4
US-08-709-209-219
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Query Match
Best Local Similarity
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Patent No. 5762938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Paolet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: PAOLETTI, Enzo
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 530 Fift
CITY: New York
STATE: NY
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
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                                                                                        LENGTH: 389 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  306
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ADDRESSEE: c/o William S. Frommer
STREET: 530 Fifth Avenue
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                                                                              linear
  96.9%;
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Score 1870; DB 1;
Pred. No. 3.4e-157;
              Length 389;
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RESULT 5
US-08-458-101-216
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Patent No. 5766599
GENERAL INFORMATION:
APPLICANT: Paoletti, Enzo
APPLICANT: Perkus, Marion E.
                                                                                                                                                  COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/458,101

FILING DATE: 01-UN-1995
INFORMATION FOR SEQ ID NO:
               CLASSIFICATION: 424
ATTORIEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 4543
TELECHOME: (212) 840-3333
TELEPHONE: (212) 840-0712
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APPLICANT: Cox, William I.

APPLICANT: Audomet, Jean-Christophe Francis
APPLICANT: Gettig, Russell Robert
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN

OF SEQUENCES: 467
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                      STREET: 530 Fifth Avenue
CITY: New York
STATE: NY
COUNTRY: USA
ZIP: 10036
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                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Curtis, Morris & Safford ADDRESSEE: c/o William S. Frommer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Johnson, Gerard P. Pincus, Steven E. Cox, William I.
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Limbach, Keith J.
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No. 5766599ton, 1
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Perkus, Marion E.
Taylor, Jill
                                                                   454310-2740
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Best Local Similarity
Matches 335; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2
Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                  APPLICANT: Audonnet, Jean-Christophe Francis
APPLICANT: Gettig, Russell Robert
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN
NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                             COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: EMP PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                186
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TYPE: amino acid
STRANDEDNESS: single
TOROTOCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      246 TTPAQQTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
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                                                                                                                                                          CITY: New York
STATE: NY
                                                                                                                          COUNTRY: U
                                                                                                                                                                                      ADDRESSEE: Curtie, Morrie & Safford ADDRESSEE: c/o William S. Frommer STREET: 530 Fifth Avenue
APPLICATION NUMBER: US/08/458,101 FILING DATE: 01-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AMOWNSTTFHOTLODPRVRGLYFPAGGSSSGAVNPVPTTASPLSSIFSRIGDPALNMENI 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AMQWNSTTFHQTLODPRVRGLYFPAGGSSSGTVNFVPTTVSPISSIFSRIGDPALKMENI 125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CPPTCPGYRMMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC
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o. 5766599
                                                                                                                                                                                                                                                                                                                    Limbach, Keit...
Limbach, Gerard P.
Johnson, Gerard P.
Pincus, Steven E.
Cox, William I
Tean-Chr/
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de Taisne, Charles
Limbach, Keith J.
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No. 5766599ton, Elizabeth
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Perkus, Marion E.
Taylor, Jill
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Pred. No. 3.4e-157;
3; Mismatches 4;
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                                                     #1.25
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TELEPAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 219:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 4543
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acids
TYPE: amino acids
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                  108 AMQWNSTTPHOTLODPRVRGLYFPAGGSSSGAVNPVPTTASPLSSIFSRÍGDPALNMENI 167
 348
               306 GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                          288 MTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                                         246 TTPAQGTSMYPSCCCTKESDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                                                                                                                                              168 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWATSLNFLGGTTVCLGQNSGSPTSNHSPTS
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                                                                                                                                                                                                                                                           AMQWNSTTFHOTLQDERVRGLYFPAGGSSSGTVNEVPTTVSBISSIFSRIGDPALMMENI 125
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Conservative
                                                                                                                                                                                                                                                                                                                                                                            96.9%; Score 1870; DB 1; 98.0%; Pred. No. 3.4e-157;
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RESULT 7 US-08-486-099-106 GENERAL INFORMATION:
APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
APPLICANT: Barney, Shawn O.
APPLICANT: Lambert, Dennis M. Sequence 106, Application US/08486099 Patent No. APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF
TITLE OF INVENTION: MEMBRANE FUSION-ASSOCIATED EVENTS, INCLUDING HEPATITIS
TITLE OF INVENTION: B VIRUS TRANSMISSION ZIP: 10036-2711

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & NUMBER OF SEQUENCES: CITY: New York STATE: New York STREET: COUNTRY: 1155 Avenue Bolognesi, Dani F Matthews, Thomas Wild, Carl T. USA Lambert, Dennis M. Dani P. Thomas J. Edmonds of the Americas Version #1.30

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RESULT 8
US-08-360-107A-116
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                                                                                                                                                                                                                                                                  Sequence 116, Application US/08360107A
PATENT NO. 6017536
GENERAL INFORMATION:
APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-486-099-106
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Best Local :
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APPLICATION NUMBER: US/08/486,099
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZİ, LAUTA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-031
TELECOMMUNICATION INFORMATION:
TELEPAN: (212) 790-9090
TELEPAN: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
TYPE: amino acid
                                                                                                                         APPLICANT: Barney, Shawn O.
APPLICANT: Lambert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION
TITLE OF INVENTION: OF MEMBRANE FUSION-ASSOCIATED EVENTS, INCLUDING HIV
TITLE OF INVENTION: TRANSMISSION
NUMBER OF SEQUENCES: 149
                                                                                                        CORRESPONDENCE ADDRESS:
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                                                  CITY: New York
                   COUNTRY:
                                                                       STREET:
                                                                                      ADDRESSEE:
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TOPOLOGY: unknown
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Local Similarity 98.2%;
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RY: USA
10036-2711
                                New York
                                                                   E: Pennie & Edmonds
1155 Avenue of the /
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                                                                                                                                                                          Sequence 106, Application US/08484223B
Patent No. 6020459
GENERAL INFORMATION:
APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Mild, Carl T.
APPLICANT: Barney, Shawn O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 96.8%;
Best Local Similarity 98.2%;
Matches 336; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/360,107A
FILING DATE: 20-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 116:
SEQUENCE CHARACTERISTICS;
LENGTH: 389 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: COTUZZÍ, LAURA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-(
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                             APPLICANT: Lambert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF
TITLE OF INVENTION: TRANSMISSION
TITLE OF INVENTION: TRANSMISSION
                    CORRESPONDENCE ADDRESS:
                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                      306 GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
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& Edmonds LLP
                                       245
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Pred. No. 5.2e-157;
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                                                                       INCLUDING HIV
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STREET: 1155 Avenue CITY: New York

of the

Americas

New York

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US-08-919-597-106
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MOLECULE TYPE:
-08-484-223B-106
                                                                                                                                  Sequence 106, Application US/08919597
Patent No. 6054265
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 106:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PATENTIN Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/484,223B PILING DATE: 07.-JUN-1995 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: CORUZZÍ, LAURA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-029
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
APPLICANT: Bolognesi, Dani P.
APPLICANT: Wild, Carl T.
APPLICANT: Wild, Carl T.
APPLICANT: Barney, Shawn O.
APPLICANT: Lambert, Dennis M.
APPLICANT: Lembert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
APPLICANT: Langlois, Alphonse J.
APPLICANT: Langlois, Alphonse J.
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Local Similarity 98.2%;
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Pred. No. 5.2e-157;
2; Mismatches 4; Indels
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TELEFAX: (214, TELEFAX: 66141 PENNIE TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: 106: SEQUENCE CHARACTERISTICS: LENGTH: 389 amino acids TYPE: amino acid
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Matches 336;
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Best Local Similarity 98.2%;
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CRASSIFICATION DATA:
PRIOR APPLICATION UNMBER: US 08/470
APPLICATION NUMBER: US 08/470
PILING DATE: 06-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAURA A.
REGISTRATION NUMBER: 30,742
REGISTRATION NUMBER: 30,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: OF TITLE OF INVENTION: TR NUMBER OF SEQUENCES: 2 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 78
TELECOMMUNICATION INFORMATION:
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STREET: 11
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228
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                                                             288
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TOPOLOGY: un
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TELEFAX: (212) 869-9741/8864
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                                                            MTTAQCTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
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                                                                                                                                                                                                                                                                   AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                             TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                                                                                         CPPTCPGYRWMCLRRPITFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC
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Pred. No. 5.2e-157;
2; Mismatches 4;
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RESULT 11
US-08-475-668A-106
; Sequence 106, Application US/08475668A
; Patent No. 6060065
; GENERAL INFORMATION:

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RESULT 12
US-08-485-551A-106
; Sequence 106, Application US/08485551A
; Patent No. 6068973
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NAMB: COTUZZI, LAURA A.

REGISTRATION NUMBER: 30,742

REFERENCE/DOCKET NUMBER: 7872-(
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090

TELEPAX: (212) 869-9741/8864

TELEPAX: (214) PENNIE
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,668A
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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APPLICANT:
APPLICANT:
                                                                                                                  348
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ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                           288 MTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
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CITY: New York
STATE: New York
COUNTRY: USA
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                                                                                                                               GLSPTVWLSVIWMMYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                          GLSPTVWLSVIWMWWWGPSLYSILSPFLPLLPIFFCLWVYI 389
                                                                                                                                                                                                TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSMAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                                                                                                                                                                                                                                   CPPTCPGYRMMCLRREIIFLFILLCLIFILVLLDYQGMLFVCFLIFGSSTTSTGFCRTC 245
                                                                                                                                                                                                                                                                                                                                            TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAGAFGLGFTPPHGGLLGWSPQAQGILQTLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAGAFGIGFTPPHGGILGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                                                                                                                                                                               CPPTCPGYRWMCLRRF1FLF1LLLCL1FLLVLLDYQGMLPVCPL1PGSSTTSTGPCRTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          389 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96.8%; Score 1868; DB 3; Length 389; larity 98.2%; Pred. No. 5.2e-157; Conservative 2; Mismatches 4; Indels
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Lambert, Dennis M.
Petteway, Stephen R.
VENTION: COMPOSITIONS FOR INHIBITION OF MEMBRANE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unknown
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CITY:
STATE: New
COUNTRY: USA
ZIP: 10036-2711

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER INM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.3
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,551A
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-023
TELEPHONE: (212) 869-9741/8864

TELEPAX: (212) 869-9741/8864

TELEPAX: SEQ ID NO: 106:

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Best Local Similarity
Matches 336; Conserv
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MOLECULE TYPE: protein
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APPLICANT: Lambert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: METHODS FOR INHIBITION OF
TITLE OF INVENTION: FUSION-ASSOCIATED EVENTS,
TITLE OF INVENTION: TRANSMISSION
                                                                                                                                                                         186
                                                                                                                                                                                                                                              126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 389 amino acids TYPE: amino acid
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STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
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                    GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                                                                                                                   TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
                                                             MTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAPGKFLWBWASARPSWLSLLVPFVQWFV
                                                                                  TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
                                                                                                                               CPPTCPGYRWMCLRRFIIFLFIILLCLIFLLVLLDYQGMLFVCFLIFGSSTTSTGFCRTC
                                                                                                                                                CPPTCPGYRWMCLRREIIFLFILLLCLIFLLVLLDYQGMLDVCFLIPGSSTTSTGPCRTC
GLSPTVWLSVIWMWYWGPSLYSILSPFLPLLPIFFCLWVYI
                                                                                                                                                                                                                                                                        AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVLTTASPLSSIFSRIGDPALNMENI
                                                                                                                                                                                                                                                                                                                                          GAGAFGLGFTPPHGGLLGWSPQAQGILQTLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 107
                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                 96.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     211
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 1868; DB 3;
Pred. No. 5.2e-157;
2; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INCLUDING
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 389;
                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 13
US-08-471-913A-106
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/471,913A
PILING DATE: 07-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAURA A.
REGISTRATION NUMBER: 30,742
REGISTRATION NUMBER: 7872-030
REFERENCE/DOCKET NUMBER: 7872-030
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 96.8%;
Best Local Similarity 98.2%;
Matches 336; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
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APPLICANT: Lambert, Dennis M.
APPLICANT: Petteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF MEMBRANE
TITLE OF INVENTION: FUSION-ASSOCIATED EVENTS, INCLUDING EPS1
TITLE OF INVENTION: TRANSMISSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Bolognesi,
APPLICANT: Matthews, '
APPLICANT: Wild, Carl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
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STATE: New Yor
COUNTRY: USA
    288
                           246 TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
                                                                                                                                                                                                                                       126 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                                                                                                                                                                                                                                      108 AMOWNSTTFHOTLODPRVRGLYFPAGGSSSGTVNPVLTTASPLSSIFSRIGDPALNMENI 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 10036-2711
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                                                                                                                                                                                                                                                                                                                 66 AMOMNSTTFHOTLODPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALMMENI 125
                                                                                                                                                                                                                                                                                                                                                                               48 GAGAFGLGFTPPHGGLLGWSPQAQGILQTLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                       6 GAGAFGLGETEPHGGLLGWSPQAQGILETLPANPPPPASTNRQSGRQPTPLSPPLRNTHPQ
WTTAOGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVOWFV 347
                                                                                              CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC 287
                                                                                                                            CPPTCPGYRWMCLRRPIIFLFILLLCLIFLLVULDYQGMLPVCPLIPGSSTTSTGPCRTC 245
                                                                                                                                                                                          TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 227
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 1868; DB 3; Length 389; Pred. No. 5.2e-157; 2; Mismatches 4; Indels
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COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARRE: PATENTIAL Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,264A
FILING DATE: 07-UNM-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LBURG A:
NAME: COTUZZI, LBURG A:
10.742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 106, Application US/08485264A Patent No. 6228983
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APPLICANT: Bologne
APPLICANT: Matthe
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (212) 869-9741/886
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
LENGTH: 389 amino acids
TYPB: amino acid
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APPLICANT: Barney, Shawn O.

APPLICANT: Lambert, Dennis M.

APPLICANT: Lambert, Stephen R.

APPLICANT: Lambert, Stephen R.

APPLICANT: Lamgiois, Alphonis FOR INHIBITION OF TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF INVENTION: MEMBRANE FUSION-ASSOCIATED EVENTS, INCLUITITLE OF INVENTION: RESPIRATORY SYNCYTIAL VIRUS TRANSMISSION NUMBER OF SEQUENCES: 232

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-021
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  348 GLSPTVWLSVIWMWWWGPSLYSILSPFLPLLPIPFCLWVYI 389
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                          186
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                                                                                                                                                                                                                                48 GAGAFGLGFTPPHGGLLGWSPQAQGILQTLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 107
                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                          65 GAGAFGLGFTPPHGGLLGWSDQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ
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                                                                                      TSGFLGPLLVLOAGFFLLTRILTIPOSLDSWWTSLNFLGGTTVCLGONSQSPTSNHSPTS 185
                                                                                                                                               AMOMNSTTFHOTLODERVRGLYPPAGGSSSGTVNPVLTTASPLSSIFSRIGDPALNMENI
                                                                                                                                                                         AMQMNSTTFHOTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALMMENI 125
                                                               TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
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Matthews, Thomas J.
Wild, Carl T.
                                                                                                                                                                                                                                                                                                                                                                                                                                              unknown
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                                                                                                                                                                                                                                                                                                                                      96.8%;
98.2%;
                                                                                                                                                                                                                                                                                                                  Score 1868; DB 4; Length 389; Pred. No. 5.2e-157; Mismatches 4; Indels
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CITY.
STATE: New_COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DCS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version #1.30
APPLICATION NUMBER: US/08/474,349A
FILING DATE: 07-UUN-1995
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 30,742
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 869-974/8864
*5141 PENNIE
***CS: 106:
***TCS: 10
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US-08-474-349A-106
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APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
APPLICANT: Wild, Carl T.
APPLICANT: Barney, Shawn O.
APPLICANT: Lambert, Dennis M.
APPLICANT: Letteway, Stephen R.
APPLICANT: Langlois, Alphonse J.
TITLE OF INVENTION: COMPOSITIONS FOR INHIBITION OF MEMBRANE
TITLE OF INVENTION: VIRUS TRANSMISSION

TITLE OF INVENTION: VIRUS TRANSMISSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
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126 TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: LLLC
STREET: New York
CITY: New York
TTATE: New York
                                                                108 AMQMNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVLTTASPLSSTFSRTGDPALNMENI 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    306 GLSPTVMLSVIMMMYMGPSLYSILSPFLPLLPIFFCLMVYI 347
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                              AMQMNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 106, Application US/08474349A
5. 6333395
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306 GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI 347
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                                                                                      186 CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC 245
                                                                                                                                           168
                                          288 MTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARPSWLSLLVPPVQWFV 347
                                                       TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKPLWBWASARPSWLSLLVPFVQWFV 305
                                                                                                                                    227
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2003, 09:27:09

Search completed: February 3, Job time: 23.1585 secs

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                1868
1803.5
1783
1786
1752
1752
                                                                                                                                                                                                                                                                   Score
                                                                                                                                                                            1929
1879
1870
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                100.0
97.4
96.9
96.9
96.8
93.5
93.5
92.4
91.0
90.8
                                                                                                                                                                                                                                                                   Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  908470 segs, 133250620 residues
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1929
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT: *
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1984.DAT: *
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1987.DAT: *
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1989.DAT: *
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1999.DAT: *
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1999.DAT: *
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1992.DAT: *
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1992.DAT: *
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1992.DAT: *
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1993.DAT: *
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1995.DAT: *
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Copyright (c) 1993 - 2003 Compugen Ltd
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              389
389
389
393
393
399
389
                                                       13
13
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22
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22
22
                       AAH10596
AAG66931
AAR27474
AAR27477
AAR27477
AAR2662931
AAR114097
AAR366932
AAR366932
AAR960794
                                                                                                                                                                                                                                                              ij
              AAG66930
                                                                                                                                                                                                                                                                                                                            SUMMARIES
Amino acid sequenc
Hepatitis B virus
HBV genotype B pre
Hepatitis B virus
Adr-type hepatitis
HBV genotype C pre
                                                                                                                                                                                                                                                       Description
                                                                                                                                   HBV genotype D pre
lpsAg protein. Sy
I3L promoter/S12/c
                                                                                                                                                                                                       HBV fusion protein
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ALIGNMENTS

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Particle for cell-specific gene delivery, useful in gene therapy,	WPI; 2000-514959/46. N-PSDB; AAA71734.	Hildt E, Hofschneider P;	(HILD/) HILDT B.	05-FEB-1999; 99DE-1004800.	04-FEB-2000; 2000WC-DE00363.	10-AUG-2000.	WO200046376-A2.		Symthetic A thice.	Hepatitis b virus	large surface protein; core antigen; gene therapy.	virus protein; cell permeability; cell-specific hinding office; rus.	Fusion protein; protein coat; virus-specific packaging signal, pri	may rusion protein comprising LHB and RGD.			AAB10596;	DI 1 0596 AAB10596 standard; Protein; 347 AA.

ИВУ

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RESULT 2
AAG66931
ID AAG6
XX AAG
AC AAG
XX HEP
XX HEP
XX HEP
XX HEP
XX HEP
XX WO2C
PN 
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Best Local S
Matches 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel particle (A), comprising a protein coat with a fusion protein (FP), and, inside the coat, a nucleic acid (I) including the sequence for a virus-specific packaging signal (psi) and a structural gene. PP contains a virus protein (VP), a peptide (P) that mediates cell permeability and a heterologous cell-specific binding site (RgD). The invention also describes (I) producing (A) in which FP (A) producing (A) in which FP (B) producing (A) in which FP (B) and (B) preparing (A) in which FP contains an HBV core antigen (HBcAg), (P) and RGD; (3) FP; (4) DNA encoding FP; and (5) expression vector containing the DNA of (d). The products of the invention are used in gene therapy of cells and tissues, in vivo or ex vivo. This sequence represents a fusion protein which is described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprises nucleic of viral protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                       19-OCT-2001
                                                                                                                                                                                                                                     Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                              HBV genotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAG66931 standard;
03-DEC-1999;
07-DEC-1999;
                                                                               20-NOV-2000; 2000WO-EP11526
                                                                                                                                                                                                                                                                                       HBeAg; antiviral;
                                                                                                                                                                                                                                                                                                               Hepatitis B virus; HBV; preCore;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    181
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les 347; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NMENITSGFLGPLLVLQAGFFLLTRILTIPQSLDSWMTSLNFLGGTTVCLGQNSQSPTSN 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HSPTSCEPTCPGYRMMCLRRFIIFLFILLLCLIFLLVLLDYQGMLFVCFLIFGSSTTSTG 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NTHPQAMOMNSTTFHQTLODDRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPAL 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14; Fig 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VOMFVGLSPTVMLSVIMMMYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCRTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCRTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPF 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HSPTSCPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTG
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ilarity 100.0%;
Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                      preS1/preS2/HBsAg
  99EP-0870252.
99US-0169287.
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                                                                                                                                                                                                                                                                                         vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein; 389
                                                                                                                                                                                                                                                                                       eCore; Core; preS1; preS2; HBS; HBX; HBPO1; genotype G; genotyping; HBcAg; HBeAg.
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Pred. No. 9.2e-143;
); Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                      polypeptide
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The invention relates to the complete nucleic acid sequence of a new CC human hepatitis B virus (HBV) genotype, provisionally named genotype CC G. This genotype was found with a high prevalence in patients CC chronically infected with HBV and residing in Burope and the USA. The CC invention relates to a fully defined sequence of 3248 nucleotides as CC given in specification, a sequence with 92% identity to the given CC sequence, or sequence that is degenerate to the mentioned sequences. CC These polynucleotides are useful for HBV genotyping. The proteins CC encoded by the polynucleotides are useful for detecting antibodies in CC encoded by the polynucleotides are useful for detecting antibodies CC directed against the proteins are useful for detecting antibodies CC and for detecting HBCAg and HBCAG (precore precursor proteins). They CC are also useful for preparing a vaccine or medicament for treating CC are also useful for preparing a vaccine or medicament for treating CC infections. The present sequence is provided in an amino acid CC sequence alignment of the preS1, preS2 and HBCAG open reading frame CC of the different HBV genotypes.
RESULT 3
AAR27472
ID AAR2
XX
AC AAR2
AC AAR2
XX
DT 24-F
XX
DE 1ps#
XX
KW Hep#
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel isolated and/or purified hepatitis B virus polypeptide and polymucleotide sequences that are phylogenetically different from genotype A-F molecules, useful for HBV diagnosis, prophylaxis and therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Fig
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                                                                                                                          AAR27472 standard;
                                                                                            AAR27472;
                              lpsAg protein.
                                                               24-FEB-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         y Match 97.4%;
Local Similarity 98.5%;
hes 337; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ
                                                                                                                                                                                                 GLSPTVMLSVIMMMYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                           CPPTCPGYRMMCLRRF11FLF1LLLCL1FLLVLLDYQGMLPVCPL1FGSSTTSTGPCRTC
                                                                                                                                                                                                                                                                                                                                                                                       TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
                                                                                                                                                                                                                                                            TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                                                                                                                                                                                                                                                                                             CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLDVCPLIPGSSTTSAGPCRTC
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                                                               (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6; 94pp; English
                                                                                                                           Protein; 389
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د.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 1879; DB 22;
Pred. No. 8.4e-139;
2; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   389;
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0

167 125 107 65

305

245 227 185

287

Hepatitis B virus;

HBV; M

protein; small

pre-S

antigen;

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The sequence given is encoded by an expression vector which comprises the hepatitis B virus (HBV) L protein (large pre-S antigen, lpsAg) gene linked to the cowpox hemorraghic region (u) promoter. This DNA sequence was used in the construction of a NYVAC recombinant construction. These were HBV genes were also used in the construction of a fusion you used in the construction. These were HBV genes were also used in the construction (small pre-S antigen, spsAg) and a fusion protein composed of the entire pre-S region (S12/core, or three distinctions of these gene sequences were inserted individually into three different sites of NYVAC separated by from each other by care the construction of vaccinia but containing essential genes. NYVAC is a copenhagen vaccine strain of vaccinia virus which has been modified by deletion of six non-essential regions of the genome encoding known or recipient loci for the insertion of foreign genes. The spacing of the three inserted sequences ensured that any recombination that did occur would lead to disruption of the vaccinia genome and would cause unviable vaccinia virus. See also AAQ35501-864.
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Matches
                                                                                                                                                              Query Match
Best Local
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11-JUN-1991;
06-MAR-1992;
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Riviere
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          126
                                     108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Fig 11; 456pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cox WI,
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                                   AMQWINSTTFHQTI
                                                                                                 GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPPASTNRQSGRQPTPLSPPLRNTHPQ
                                            AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI
                                                                                ĠĀĠĀPĠĹĠ₽ŦPPHĠĠĹĹĠŴŚPQĀQĠĬĹQŦĹPĀNPPPĀSTNRQTĠŔQ₽ŦPĹŚPPĹRNTHPQ
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                                                                                                                                                              Similarity
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K, Paoletti B,
J, Taylor J;
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3, Perkus MB,
                                                                                                                                                        DB 13;
1.2e-138;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pox-virus - use for as rabies, hepatitis
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3, Pincus SE;
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The sequence given is encoded by an expression cassette which comprises the hepatitis B virus (HBV) L protein (large pre-S antigen, lpsAg) gene which is precisely linked to the Amsacta moorei entomopoxvirus (AmBPV) 42 kD promoter. This DNA sequence was used in the construction of a NYVAC recombinant expressing the HBV gene. Other HBV genes were also used in the construction. These were HBV M

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RESULT 4
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11-JUN-1991;
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Limbach
                                                                                                                                            Disclosure;
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91US-0713967.
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K, Paoletti
J, Taylor J
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B, Perkus N
                                                                                                                                                                                                pox-virus -
as rabies, l
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    use for hepatitis

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                                                   (INRM ) INSERM INST NAT SANTE & (INWR ) BIOMERIEUX SA.
                                                                                                                                                                                                                                                                                                                                                         Hepatitis B virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV 305
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98.0%;
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pred. No. 4.2e-138;
3; Mismatches 4;
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           Chemin I,
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              Komurian-Pradel
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RESULT 6
AAU1407
ID AAU1407
XX AAU1407
XX AAU1407
XX Hepa
XX Hepa
XX Anti
KW Majc
XX Mart
XX Hepa
XX Anti
XX Mart
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents a mutated Pre-S gene. It is part of the mutant hepatitis B virus (mHVB) of the invention, which has a circular, partially double-stranded genome comprising pre-S, S, C, P and X genes, encoding, respectively, surface antigens, envelope protein (HBsAg), proteins HBeAg and HBcAg, DNA polymerase/reverse transcriptase, and HBsAg. The mutant HBV of the invention are used for induction of specific humoral or cellular immune responses, or antisense inhibition of viral protein expression. Proteins and nucleic acids derived from mHVB (also related antibodies, vectors, transformed cells and coligonucleotides) are useful for diagnosis (in standard amplification, hybridization or immune-complex forming assays) and for treatment of, or vaccination against, HBV infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatitis B virus major surface antigen precursor
                                                                                                                                                                                                                                                                                                                                                       21-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU14097 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             168
                                                                                                                                                                                                                      major
                                                                                                                                                                                                                                            Anti-retroviral; DP178-like; DP107-like; antifusogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        186
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                                                                                                                                                             Hepatitis B virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GLSPTVMLSVIMMMWYMGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TSGFLGPLLVLQAGEFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CPPTCPGYRMMCLRRFIIFLFILLLCLIFLLVLLDYQGMLFVCFLIFGSSTTSTGFCRTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWTTSLNFLGGTTVCLGQNSGSPTSNHSPTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CPPTCPGYRWMCLRRFIIFLFILLICLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
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                                                                                                                                                                                                                   surface antigen precursor S; HIV
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                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                   subtype AYW.
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Pred. No. 6.1e-138;
2; Mismatches 4;
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                                                                                                                                                                                                                                                                antiviral;
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05-JUL-2000; 2000WO-US35727

19-JUL-2001

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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to peptides which exhibit anti-retroviral cactivity. The peptides of the invention (AAU12559-AAU14009) comprise CD pp178-like and Dp107-like peptides. The Dp178 peptide corresponds CC corresponds 639-673 of the transmembrane protein gp41 from human CC corresponds to amino acids 558-595 of gp41 from HIV-1LAI. The invention CC corresponds to amethod of identifying compounds that inhibit the CC also relates to a method of identifying compounds that inhibit the CC detecting the formation of a Dp107/Dp178 complex. The method comprises CC or absence of a test compound, in a reaction mixture containing Dp107 cand Dp178 peptides. The method is useful for identifying compounds, CC and Dp178 peptides. The method is useful for identifying compounds, CC antifusogenic, antiviral or intracellular modulatory activity. The CC pp178-like/Dp107-like peptides are useful to inhibit human and non-human CC retroviral, particularly HIV, transmission to uninfected cells. The CC present sequence represents a peptide sequence from Hepatitis B virus extraction may make the sequence represents a peptide sequence from Hepatitis B virus extraction for the sequence constant of the sequence constant of the sequence represents a peptide sequence from Hepatitis B virus extraction for the sequence constant of the sequence constant of the sequence constant of the sequence constant of the sequence from Hepatitis B virus extraction for the sequence constant of the sequence consta
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                19-OCT-2001
                                                            AAG66932;
                                                                                                 AAG66932 standard; Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying a compound that inhibits the formation of or disrupts a DP107/DP178 complex, especially compounds with antifusogenic, antiviral or intracellular modulatory activity, by detecting the formation of a DP107/DP178 complex -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   169
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                                                                                                                                                                                                     WFVGLSPTVWLSVIWMWWWGPSLYSILSPFLPLLPIFFCLWVYI 393
                                                                                                                                                                                                                               WEVGLSPTVWLSVIWMMWYWGPSLYSILSPELPLLPIEFCLWVYI 347
                                                                                                                                                                                                                                                                                                                  RTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQ 302
                                                                                                                                                                                                                                                                                                                                                                                               PTSCPPTCPGYRWMCLRRFIIPLPILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                enitsgflgfllvloagffiltriltifosldswatslnflggttvclgonsosftsnas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENITSGELGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PQAMQMNSTTF-HQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALIM 122
                                                                                                                                                                                                                                                                                       RTCMTTAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAPGKPLWEWASARFSWLSLLVPFVQ
                                                                                                                                                                                                                                                                                                                                                                          PTŚCEPTCPĠYRWYCLKRPIIPLPILLLCLIPLLVLLDYQGWLPVCPLIPGŚŚTTŚTĠPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PQAMQMNSTTFIIQTLQDPRVRGLYFPAGGSSSGTVNPVLTTASPLSSIFSRIGDPALNM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAGAFGLGFTPPIIGGLLGWSPQAQGILQTLPANPPPASTNRQSGRQPTPLSPPLRNTII 108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lackey JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      393 AA;
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             (first entry)
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                                                                                                 399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 1803.5;
Pred. No. 6.7e.
2; Mismatches
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3.7e-133;
7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to the complete nucleic acid sequence of a new CC human hepatitis B virus (HBV) genotype, provisionally named genotype CC of This genotype was found with a high prevalence in patients of the USA. The cohornically infected with HBV and residing in Europe and the USA. The CC invention relates to a fully defined sequence of 3248 nucleotides as CC given in specification, a sequence with 92% identity to the given ce that is degenerate to the mentioned sequences. These polynucleotides are useful for HBV genotyping. The proteins CC encoded by the polynucleotides are useful for detecting antibodies in CC a biological sample. Ligands that bind to the proteins and antibodies are of intected against the proteins are useful for detecting the proteins CC and for detecting HBCAg and HBCAG (precore precursor proteins). They are also useful for preparing a vaccine or medicament for treating CC sequence alignment of the pressl, pres2 and HBCAG open reading frame CC of the different HBV genotypes.
                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                        Matches 318;
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated and/or purified hepatitis B virus polypeptide a polynucleotide sequences that are phylogenetically different f genotype A-F molecules, useful for HBV diagnosis, prophylaxis
                                                                            238
                                                                                                                                          178
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                                                                                                                                                                          126
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07-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis B virus; HBV; preCore; Core; preS1; preS2; HBS; HBX; HBPol;
HBsAg; antiviral; vaccine; genotype G; genotyping; HBcAg; HBeAg.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HBV genotype E preS1/preS2/HBsAg
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MTLAQGTSNFPSCCCSKPSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFA
                    TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKELWEWASARFSWLSLLVDEVQWEV
                                                                CPPICPGYRWMCLRRFIIFLFIIILLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC
                                                                                     CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVILLDYQQMLPVCPLIPGSSTTSTGPCRTC
                                                                                                                                                       TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS
                                                                                                                                                                                                                                                                                    GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                                                                                                                                TŚGFLGPLLVLQAGFFLLTKILTIPQSLDSWWTSLNFLGGAPVCLGQNSQSPTSNHSPTS
                                                                                                                                                                                               AMQWNSTTFHQALQDPRVRGLYFPAGGSSSGTVNPVPTTASLISSIFSRIGDPAPNMESI
                                                                                                                                                                                                                       AMQWNSTTFHQTLQDPRVRGLYPPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI
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99US-0169287.
                                                                                                                                                                                                                                                                                                                                 92.4%; Score 1783; DB 22;
93.0%; Pred. No. 2.7e-131;
cive 11; Mismatches 13;
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GLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFPCLWVYI

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358

GLSPTVMLSVIMMMWMMGPSLYDISTPIPLLPIPFCLMVXI

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RESULT 8
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Matches 312
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05-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-OCT-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hepatitis C virus; HCV; hepatitis B virus; HBV; immunisation; vaccine; infection; core protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis B virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A new fusion protein comprises a hepatitis B virus (HBV) S gene protein (AA736611) coupled to amino acids 1-69, 1-70 or 1-154 of the hepatitis C virus (HCV) core protein (AA728348).

The nucleic acid has the coding sequence linked to a CMV promoter, RSV enhancer, polyadenylation sequence and opt. the S'-UTR of HCV.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid encoding hepatitis B and C virus fusion proteins -incomplete hepatitis C virus genome, are useful in vaccines for prevention or treatment of HBV and HCV infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-209642/21.
N-PSDB; AAT36611.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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(GEHO) GEN HOSPITAL CORP
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                                                                                                                                                                 AMQWNSTTFHQTLQDBRVRGLYFPAGGSSSGTVNPVPTTVSDISSIFSRIGDPALMMENI 125
                                                                      CPPICPGYRMMCLRRFIIFLFILLCLIFLLVLLDYOGMLPVCPLIPGSSTTSTGPCRTC CPPICPGYRMMCLRRFIIFLFILLCLIFLLVLLDYOGMLPVCPLLPGTSTTSTGPCKTC
                                                                                                                                                                                                                                                              TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAPGKFLWEWASARFSWLSLLVPFVQWFV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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ilarity 91.2%;
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94US-0318248
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Pred. No. 3.5e-129;
.1; Mismatches 19;
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RESULT 9
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Best Local Similarity 90.9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Adr-type hepatitis B virus SA,
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N-PSDB; AAN60714.
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                                                                                                                                                                                                                                                                                                                                                                                   The sequence allows for the preparation of adr-type hepatitis B surantigens for vaccine, without the use of human blood. The proteins expressed from human hepatoma cell strain huGK-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADR-type hepatitis-B surface antigen f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NICA+) JAPAN FOUND CANCER (MEIP) MEIJI MILK PROD CO (GANK+) GAN KENKYU KAI ZH.
                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Fig 5; 30pp; Japanese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GLSPTVWLSVIWMMYWGPSLYNILSPFLPLLPIFFCLWVYI
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                             TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSPTSNHSPTS 185
                                                                                 AMOWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                                                                       GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ
  TSGFLGPLLVLQAGEFLLTRILTIPQSLDSWWTSLNFLGGAPTCPGQNSQSFTSNHSPTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUGANO H,
                                                                                                                                                                                                                                                                                                                                                    389 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /label= 31K-36K intermediate surface
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                                                                                                                                                                                                                                                                  Score 1752; DB 7
Pred. No. 7e-129;
L2; Mismatches 1
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n hugk-14 avoiding use of human blood
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RESULT 10
AAG66330
AD AAG66
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XX
                                                                                                                                                       CC The invention relates to the complete nucleic acid sequence of a new CC G. This genotype was found with a high prevalence in patients CC chronically infected with HBV and residing in Europe and the USA. The CC circunically infected with HBV and residing in Europe and the USA. The CC given in specification, a sequence with 92% identity to the given CC sequence, or sequence that is degenerate to the mentioned sequences. CC These polynucleotides are useful for HBV genotyping. The proteins CC encoded by the polynucleotides are useful for detecting antibodies in CC abiological sample. Ligands that bind to the proteins and antibodies and for detecting HBcAg and HBcAg (precore procursor proteins). They can detecting HBcAg and HBcAg (precore procursor proteins). They CC are also useful for preparing a vaccine or medicament for treating CC are also useful for preparing a vaccine or medicament for treating CC are also useful for present sequence is provided in an amino acid CC sequence alignment of the preS1, preS2 and HBcAg open reading frame CC of the different HBV genotypes.
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                                                                                                                                   Novel isolated and/or purified hepatitis B virus polypeptide and polynuclectide sequences that are phylogenetically different from genotype A-F molecules, useful for HBV diagnosis, prophylaxis and
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HBsAg; antiviral; vaccine; genotype G; genotyping; HBcAg; HBeAg.
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The invention relates to the complete nucleic acid sequence of a new human hepartitis B virus (HBV) genotype, provisionally named genotype G. This genotype was found with a high prevalence in patients chronically infected with HBV and residing in Europe and the USA. The invention relates to a fully defined sequence of 3248 nucleotides as given in specification, a sequence with 92% identity to the given gequence, or sequence that is degenerate to the mentioned sequences. These polymucleotides are useful for HBV genotyping. The proceins encoded by the polymucleotides are useful for detecting antibodies in a biological sample. Ligands that bind to the proteins and antibodies

antibodies in and antibodies

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Claim 16;

Fig

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                                                                                                                                                                                                                                                    Egg white lysozyme; hepatitis B virus (HBV) surface antigen(HBsAg); Saccharomyces cerevisiae AH22R/pGLD LP39-RcT; L protein; M protein; Saccharomyces cerevisiae LP31-
                                                                                                                                                                                                                                                                                              Sequence of adr type (HBsAg) L protein (M
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Peptide(s) having hepatitis B surface antigenicity - used as vaccine for prevention of hepatitis B virus and in diagnostic kits
                                                                                                                                                                                                                   Hepatitis B
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                                                                                                                                                   13-APR-1988;
                                                                                                                                                                        26-OCT-1988
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                                                             WPI; 1988-301233/43
                                                                                 Fujisawa
                                                                                                                              12-OCT-1987;
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pred. No. 3.7e-128;
l3; Mismatches 20;
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CC CINTENDUCED IN 18 AN ENV protein. When L protein genes are CC introduced into animal cells (CHO cells) they produce HBSAg particles CC containing M and S proteins. The advantage is that peptides having CC HBSAg antigenicity are excreted outside the cells and easily purified. CC Claimed is a eukaryotic cell transformed with the rDNA. Pref. CC The cell is a yeast cell, esp. S. cerevisiae AH22R-/pGLD LP39-CC THE CELL, and is bound to the 5'-terminal of a DNA coding for a signal peptide which functions in a eukaryotic CC cell, and is bound to the 5'-terminal of a DNA coding for a peptide CC cell, and is bound to the 5'-terminal of a DNA coding for a peptide CC cell, and is bound to the 5'-terminal of a DNA coding for a peptide CC cell, and is bound to the 5'-terminal of a DNA coding for a peptide CC cell, and is bound to the 5'-terminal of a DNA coding for a peptide CC cell, and is bound to the 5'-terminal of a DNA coding for a peptide of CC cell, and is bound to the 5'-terminal of a peptide of CC cell, and is bound to the 5'-terminal of a signal peptide of CC cell, and is bound to the signal peptide may be a signal peptide of CC cell, and is bound to the signal peptide of CC cell, and is bound to the signal peptide of CC cell, and is desirable that the genes are altered CC by the protease; therefore it is desirable that the genes are altered CC peptide containing the residue (preferably the 44-49 peptide) may be CC deleted. Preferable genes include L protein gene (altered) coding for the amino acid sequence 1-383 shown in AAN80973.
                                                                                                                                                                                                                                                                                                                        RESULT 13
AAR55287
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Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                  Surface antigen; recombinant replicable vaccinia virus; prevention; therapy; epitope; hepatitis B virus.
                                                                                                                                                                                                        Deduced
                                                                                                                                                                                                                                                                                                           AAR55287 standard; Protein; 389
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                                                                                                                 Hepatitis B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TIPAQGTSMFPSCCCTKPSDGNCTCIPIPSSWAFARFLWEWASVKESWLSLLVPFVQWFV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTPAQGTSMYDSCCCTKDSDGNCTCIPIPSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
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                                                                                                                                                                                                      sequence of HBV surface antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  383 AA;
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                                                                                                                       virus
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                                                                                                                                                                                                                                          entry)
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Best Local Simi
Matches 305;
           Hepatitis B virus
                                            DNA polymerase; large surfa
transactivating X protein;
                                                                 HBV; HBV surface antigen-'S'-133 Oon strain (Met to Thr);
                                                                                        Amino acid seqeunce of a HBV large surface antigen protein
                                                                                                                             27-MAR-2000
                                                                                                                                                                         AAYS4045 standard; Protein;
                                                                                                                                                      AAY54045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The HBV genome (subtype adw) is contained in plasmid pAM6. The 3' end of the ORF conding for the three surface Ags was manipulated to generate pRO-02 which contains an intact ORF capable of encoding LS, MS and S Ags. The DNA sequences encoding S, MS and LS were subcloned and sequenced. The sequences of the LS are AAQ67970/R55287; the sequences of MS are AAQ67961/R55281 and The sequences of S are AAQ67962/R55282. Changes in nucleotide and deduced AA sequences from the published sequence are believed to result from variability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prevention and treatment of hepatitis - using recombinant replicable vaccinia viruses contg. hepatitis B virus surface and
                                                                                                                                                                                                                                                 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                          GLSPTVWLSAIWAWWYWGPSLYSIVSPFIPLLPIPFCLWVYI
                                                                                                                                                                                                                                                       GLSPTVWLSVIWMMYWGPSLYSILSPFLPLLPIFFCLWVYI 347
                                                                                                                                                                                                                                                                                                          TTPAQGTSMYPSCCCTKPSDGNCTCIPIDSSWAFGKFLWEWASARFSWLSLLVPFVQWFV
                                                                                                                                                                                                                                                                                                                                                              CPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSSTTSTGPCRTC
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                                                                                                                                                                                                                                                                                                                                             ĊPPIĆPGYRMMĊLRRPIIPLFILLLĆLIPLLVLLDYOGMLPVČPLIPGSTTTSTGPĆKTĆ
                                                                                                                                                                                                                                                                                                                                                                                                TSGPLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGSPVCLGQNSQSPTSNHSPTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GVGAPGPGPTPPHCGVLGWSPQAQGILTTVSTIPPPASTNRQSGRQPTPISPPLRDSHPQ 107
                                                                                                                                                                                                                                                                                           TTPAQGNSMPPSCCCTKPTDGNCTCTPIPSSWAPAKYLWEWASVRPSWLSLLVPFVQWFV
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                                                                                                                           (first
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                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            89.5%; Score 1727; DB 15; 89.2%; Pred. No. 6.3e-127; tive 14; Mismatches 23;
                                         surface antigen; core protein;
tein; hepatitis vaccine; HBV in
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                                                                                                                                                                                                                                                                                                                           The present sequence is encoded by the genome of an isolated strain of C Hepatitis B virus designated human Hepatitis B virus (HBV) surface C antigen-'S'-133 Oon strain (Met to Thr). The viral genome is deposited C as ECC accession numbers psy121501, psy121502 and psy121503. The C uncleotide sequence was isolated from hepatocellular carcinoma (HCC). C The nucleotide sequence encodes four overlapping proteins, which are C uncleotide sequence encodes four overlapping proteins, and a C transactivating X protein. The large surface antigen, a core protein, and a C wild type sequence in that it contains a Thr at position 133 of the C wild type sequence instead of a Met. The proteins are used to produce C antibodies. The proteins, polynucleotide and antibodies can be used for C detecting the novel HBV strain. The HBV polypeptides can be used C in hepatitis vaccines. The HBV novel strain polypeptides can be used to identify compounds for treating or preventing HBV infection or chematocellular carcinoma
                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                      Matches 308;
                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 23; Page 39-40; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated hepatitis B virus strain, useful for, e.g. treatment hepatitis infection - \ensuremath{\mathsf{P}}
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                                                   GAGAFGLGFTPPHCGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
                TTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKF1WEWASARFSWLSLLVDPVQWPV
                                                                                                                      TSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTVCLGQNSQSFTSNHSPTS
TTPAQGNSTFPSCCCTKPSDGNCTCIPIPSSWAFARPLWEWASVRFSWLSLLVPFVQWFV
                                                                                                     TSGFLGPLLVLQAGFESLTRILTIPQSLDSWWTSLNFLGGAPTCPGQNSQSPTSNHSPTS
                                                                                                                                                         ATOWNSTTFHOALLDPRVRGLYFPAGGSSSGTVNPVPTTASPISSIFSRTGDPAPNMENT
                                                                                                                                                                        AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000-106104/09.
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                                                                                                                                                                                                                                                                                                                          400 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zhao Y,
                                                                                                                                                                                                                                                                 89.2%; Score 1721; DB 21; 90.1%; Pred. No. 1.9e-126; tive 10; Mismatches 24;
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GLSPTVWLSVIWMWYWGPSLYSILSPFLPLLPIFFCLWVYI

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RESULT 15
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Best Local (
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Region
                                                                                                                                                       The modified L protein has the Gly 13 amino acid in the pres1 region of the wild-type sequence deleted. The deletion results in the synthesis of a non-myristilated L protein. (The first amino of the ad sequence are not given). The modified product may be used alone or in a composite particle comprising at least two polypeptides corresp, to all or part of a protein having the biological activity of one of the hepatitis B surface antigens. The composite particles are useful for preparing improved hepatitis B vaccines or for treating of HBV. See also AAQ10677-79.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Region
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                                                                                                                                                                                                                                                                                                               Modified hepatitis B virus large surface protein - has amino acid sequence encoding L protein used in vaccine for treating or preventing hepatitis B without effects
                                                                                                                                                                                                                                                                                                                                                                                                Comberbach M, Harford N,
Jacobe E, Hollenberg CP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                               WPI; 1991-059585/09.
N-PSDB; AAQ10678.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-AUG-1989;
                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                          Disclosure; Page 4-7; 79pp; English.
*.66 AMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIGDPALNMENI 125
                                                                                                                                                                                                                                                                                                                                                                                                                                     (SMIK ) SMITHKLINE BIOLOGIC.
                                                                                      Local Similarity 88.3
nee 302; Conservative
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                                                    GAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPPASTNRQSGRQPTPLSPPLRNTHPQ 65
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Viruses; Retroid viruses; Hepadnaviridae;
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Borisova G.P., Pumpen P.P., Bychko V.V., Pushko P.M., I

Borisova V., Gren E.Y., Tsibinogin V.V., Kukain R.A.;

Dokl. Biochem. 279:386-390(1985).
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MEDLINE=81012091; PubMed=399327;
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Submitted (AUG-1994)
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InterPro; IPR002006; Hepatitis_core.
Pfam; PF00906; Hepatitis_core; 1.
SEQUENCE 212 AA; 24348 MW; 6E910
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  PSPRRRRSQSPRRRRSQSRESQC
                                                                                  LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPI
                                                                                                       LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT 192
                                                                                                                                                                        EHCSPHHTALROAILCWGDLMTLATWVGVNLEDP---
                                                                                                                                                                                                                                                        PTVQASKLCLGWLWGMDIDPYKBFGATVELLSFLPSDFFPSVRDLLDTASALYRBALESP
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ne EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Score 1037.5; ; Pred. No. 3.3e. 1; Mismatches
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Last annotation update)
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212
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Q89597;
01-NOV-1996
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SPECIES-Hepatitis B virus; STRAIN=143-FAM G;

Zampino R., Lobello S., Chiaramonte M., Venturi-Pasini C.,

Thursz M., Karayiannis P.;

Intra-familial transmission of Hepatitis B virus in Italy:
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01-NOV-1996 (TREMBLrel. 01, Createq, 01-NOV-1996 (TREMBLrel. 10, Last sequence update) 01-NOV-1996 (TREMBLrel. 19, Last sannotation update) 01-DEC-2001 (TREMBLrel. 19, Last annotation update) HBCAG (PRE-CORE protein) (PRECORE) core protein).
                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; X72702; CAA51257.1; -.
EMBL; X80925; CAA56887.1; -.
EMBL; AF419525; AAL15953.1; -.
EMBL; AF419527; AAL15953.1; -.
EMBL; AF350127; AAK57244.1; -.
EMBL; AF350127; AAK57322.1; -.
InterPro; IPR002006; Hepatitis_core.
Pfam; PF00906; Hepatitis_core; 1.
CHAIN 30 212 CORE PROTEIN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Phylogenetic sequence analysis and amino acid variation of the core gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SPECIES-Hepatitis B virus (subtype ayw); STRAIN=AYW, Karayiannis P.;
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SPECIES=Hepatitis B virus (subtype ayw);
MEDLINE=94079539; PubMed=8257295;
Preisler-Adams S., Schlayer M.J., Peters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis B virus (subtype Hepatitis B virus. Viruses; Retroid viruses; H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
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LWEHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT
                                                                                           EHCSPHHTALRQAILCWGELMTLATWVGVNLEDP----ASRDLVVSYVNTNMGLKFRQL
                                                                                                                            EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQL
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                                                                                                                                                                                                                                                                                                                                                                                                212
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                                                                                                                                                                                                                                                                                                                 90.4%;
                                                                                                                                                                                                                                                                                                                                                                                             24336 MW;
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                                                                                                                                                                                                                                                                                             Score 1037.5;
Pred. No. 3.3e
1; Mismatches
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                                                                                                                                                                                                                                                                                                                    5; DB 12; Length .3e-97;
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193 190

PSPRRRRSQSPRRRRSQSREPQC

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Viruses; Retroid viruses; Hepadnaviridae;
NCBI_TaxID=10407;
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Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases.
EMBL; 85293; CAA59616.1; -.
Interpro; IPR002006; Hepatitis core.
Pfam; PF00966; Hepatitis core; 1.
SEQUENCE 212 AA; 24363 MW; 6940D04E90A410C3 CRC64;
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Viruses, Retroid viruses, Hepadnaviridae,
NCBI_TaxID=10407;
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PRE-C/CORE.
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PRE-C/CORE.
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                                                                                                                                                          Lai M.E., Mazzoleni A.P., Porru A., Balestrieri A.;
Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases.
EMBL; X85315, CAA596641;
InterPro, IPR002006; Hepatitis core.
Pfam; PF00906; Hepatitis core; 1.
SEQUENCE 212 AA; 24366 MW; CEABDD4E8B11E5E8 CRC64;
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Pred. No. 5.3e
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                                                       Score 1035.5; DB 12; Length 212; Pred. No. 5.3e-97;
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         Pre-c/core protein.
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Hepatitis B virus.
Viruses; Retroid viruses; F
NCBL TaxID=10407;
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SEQUENCE FROM N.A.
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Best Local S
Matches 195
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Best Local
    Q91SY7
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Pfam; PP00906; Hepatitis core; 1
SEQUENCE 212 AA; 24376 MW; D3017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.

Lai M.B., Mazzoleni A.P., Porru A., Balestrieri A.,

Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases.

EMBL; X85316; CAA59666.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis B virus.
Viruses, Retroid viruses,
NCBI_TaxID=10407,
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01-NOV-1996 (TremBLrel.
01-DEC-2001 (TremBLrel.
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01-NOV-1996
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PRE-C/CORE.
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                                                                                                      PSPRRRRSQSPRRRRSQSRESQC
                                                                                                                              PSPRRRRSQSPRRRRSQSREPQC 215
                                                                                                                                                                               LWPHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILLTLPETTVVRRRGRSPRRRT
                                                                                                                                                                                                          LWFHISCLTFGRETVIEYLVSEGVWIRTPPAYRPDNAPILSTLPETTVVRRRGRSPRRRT 192
                                                                                                                                                                                                                                                                    BHCSPHHTALRQAILCWGELMTLATWVGVNLEDP---
                                                                                                                                                                                                                                                                                        EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQL 132
                                                                                                                                                                                                                                                                                                                                                                        PTVQASKLCLGWLWGMDIDPYKEPGATVELLSFLFSDFFFSVRDILDTASALYREALESP
                                                                                                                                                                                                                                                                                                                                                  PTVQASKLCLGWLWGNDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP
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; Pred. No. 6.7e. 
0; Mismatches
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Last annotation update)
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PRT;
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Pred. No. 6.7
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212
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                                                                                                                                                                                                                                                               -- ASRDLVVSYVNTNMGLKFRQL
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EMBL; AF419526; AALL15954.1; -.

InterPro; IPR002006; Hepatitis_core.

Pfam; PF00906; Hepatitis_core; 1.

CHAIN

30 212 CORE PROTEIN.

CHAIN 30 212 CORE PROTEIN.

SEQUENCE 212 AA; 24421 MW; B76FFE275B047AD1 CRC64;
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01-DEC-2001 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
                                                                                                                                                                                                                                      STRAIN=144-FAM G;
Zampino R., Lobello S., Chiaran
Thursz M., Karaylannis P.;
Thura-familial transmission of
Phylogenetic sequence analysis
gene.";
                                                                                                                                                                                                                                                                                                                                                                                                  Pre-core protein.
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae;
NCBI_TaxID=10407;
                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q917I7
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Submitted (FEB-2001) to the EMBL/GenIEMBL; AF35015; AAK57272.1; -.

InterPro; IPR002006; Hepatitis core.

Pfam; PF00906; Hepatitis core; 1.

SEQUENCE 212 AA; 24363 MW; 27E96;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=GAM1211;
Dumpis U., Mendy M., Kai
"Prevalence of HBV core
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Precore/core protein.
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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01-DEC-2001 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
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         Conservative
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                         89.9%;
96.1%;
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96.1%; Pred. No. 1.1e
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re promoter/precore/core
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20,
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19,
20,
Score 1032.5; DB 12
Pred. No. 1.1e-96;
1; Mismatches 2;
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Chronic Carriers.";

Submitted (FEB-2001) to the EM EMBL; AF350218; AAK57335 1; ---

EMBL; AF350218; AAK57216 1;

EMBL; AF350100; AAK57221 1;

EMBL; AF350100; AAK57220 1;

EMBL; AF350100; AAK57223 1;

EMBL; AF350106; AAK57233 1;

EMBL; AF350108; AAK57233 1;

EMBL; AF350110; AAK57235 1;

EMBL; AF350110; AAK57236 1;

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EMBL; AF350110; AAK57236 1;

EMBL; AF350121; AAK57236 1;

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EMBL; AF350121; AAK57236 1;
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Hepatitis B virus.
Viruses, Retroid viruses; Hepadnaviridae;
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01-MAR-2002
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Dumpis U., Mendy M., Ka
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Viruses; Retroid viruses;
NCBI_TaxID=10407;
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SEQUENCE FROM N.A.
Lai M.B., Mazzoleni A.P., Balestrieri A., Melis A., Porru A.;
Isai M.B., Mazzoleni A.P., Balestrieri A., Melis A., Porru A.;
"Sequence analysis of HBV genomes isolated from patients with
negative chronic liver disease.";
Submitted (MAR-1992) to the EMBL/GenBank/DDBJ databases.
EMBL; X65257; CAA46350.1; -.
InterPro; IPR002006; Hepatitis core.
Pfam; PF00906; Hepatitis core; 1.
SEQUENCE 212 AA; 24348 MW; 1861AB0DA21E49E1 CRC64;
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L; AF350181; AAK57308.1; -...
L; AF350186; AAK57303.1; -...
L; AF350186; AAK57303.1; -...
L; AF350189; AAK57307.1; -...
L; AF350199; AAK57307.1; -...
L; AF350194; AAK57310.1; -...
L; AF350195; AAK57311.1; -...
L; AF350196; AAK57312.1; -...
L; AF350197; AAK57312.1; -...
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L; AF350199; AAK57314.1; -...
L; AF350210; AAK57317.1; -...
L; AF350211; AAK57317.1; -...
L; AF350212; AAK57339.1; -...
L; AF350214; AAK57339.1; -...
L; AF350214; AAK57339.1; -...
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GenCore version 5.1.3 Compugen Ltd.

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Title: Perfect score: Sequence:

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters: 112892 seqs, 41476328 residues 112892

Searched:

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : SwissProt_40:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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                                              MEDLINE-85204397; PubMed=3996597; Bichko V., Pushko P., Dreilina D., "Subtype ayw variant of hepatitis B analysis.";
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                                                                                                                                                                "Nucleotide sequence of the hepatitis cloned in E. coli.";
                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=81012091; PubMed=399327;
Galibert F., Mandart B., Fitoussi
                                                                                                                                                                                                                                                                               Viruses; Retroid
NCBI_TaxID=10418;
                                                                                                                                                                                                                                                                                                     Hepatitis B virus (subtype ayw)
Viruses; Retroid viruses; Hepac
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Matches 182;
                                                                                                     This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Enthe European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by antentities requires a license agreement (See http://www.isb-rorsend an email to license@isb-sib.ch).
                               EMBL; J02202; AAA45486.1; -. EMBL; A08967; CAA00816.1; -. PIR; B93217; NKVLA2.
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P03147;
                                                                                                                                                                                                                                                                                                 Pasek M., Goto T., Gilbert W., Zink B., Schall Leadbetter G., Murray K.; "Hepatitis B virus genes and their expression Nature 282:575-579(1979).
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MEDLINE=81012115; PubMed=399329;
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01-AUG-1990
                                                                                                                           InterPro; IPR002006; Hepatitis core. Pfam; PF00906; Hepatitis core; 1. Core protein; Repeat. REPEAT 162 169
                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                       Core antigen.
Hepatitis B virus (subtype adw / strain Japan/pJDW233).
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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                                                                                                                                                                          EMBL; D00329; -; NOT_ANNOTATED_CDS.
PIR; A28925; NKVLJ1
                                                                                                                                                                                                                                                                                                                 J. Gen. Virol. 69:2575-2583(1988).
                                                                                                                                                                                                                                                                                                                       "Typing hepatitis B virus by homology in nucleotide sequence: comparison of surface antigen subtypes.";
                                                                                                                                                                                                                                                                                                                                                       Okamoto H., Tsuda F.,
Miyakawa Y., Mayumi M.
                                                                                                                                                                                                                                                                                                                                                                MEDLINE=89010694; PubMed=3171552;
Okamoto H., Tsuda F., Sakugawa H.,
                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HPBVJ
                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10413;
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Core protein; Repeat.
REPEAT 162 169
REPEAT 170 177
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              MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL
MDÍDÞÝKEFGATVELLSFLÞSDFFÞSVRDLLDTVSALÝREALKSPEHCSPHHTALRQAIL
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                                                    Similarity 93.175; Conservative
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                                                                                                   162
170
183 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              (Rel. 15, Created)
(Rel. 15, Last sequence update)
(Rel. 17, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 183
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                                                                                                                177
                                                                                                 21224 MW; 9FDD6B5F5AF5B160 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21042 MW;
                                                 80.1%; Score 919.5; DB 1
93.1%; Pred. No. 2.3e~75;
Live 3; Mismatches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          81.2%;
93.6%;
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Pred. No. 1.6
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                                                                          DB 1;
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                                                                          183;
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                                                 5.
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                                              Gaps
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CWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQLLWFHISCLTFGRETV 147

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RESULT
CORA_H
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RESULT. 8
CORA_HPBVW .
ID __CORA_HPBVW
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Best Local !
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Hepatitis B virus (subtype adw / strain Okinawa/pODW282).
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Okamoto H., Tsuda P., Sakugawa H., Sastrosoewignjo R.I., Imai Miyakawa Y., Mayumi M.;
"Typing hepatitis B virus by homology in nucleotide sequence: comparison of surface antigen subtypes.";
J. Gen. Virol. 69:2575-2583(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=89010694; PubMed=3171552; Okamoto H., Tsuda F., Sakugawa H.,
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InterPro; IPR002006; Hepatitis_core.
Pfam; PF00906; Hepatitis_core; 1.
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REPEAT 162 10
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                                                                                                                             SQSRESQC
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(Rel.
(Rel.
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          STANDARD;
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93.1%;
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Last annotation updat
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3; Mi
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Pred. No. 2.3
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RESULT 9
CORA_HPBV4
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Best Local :
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21-JUL-1986 (Rel. (
21-JUL-1986 (Rel. (
01-OCT-1989 (Rel. )
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P03150; P03151;
21-JUL-1986 (Rel. (
21-JUL-1986 (Rel. (
01-FEB-1991 (Rel. )
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Core protein; Repeat.
REPEAT 164 171
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MEDLINE=83168919; PubMed=6300776;

MEDLINE=83168919; PubMed=6300776;

Ono Y., Onda H., Sasada R., Igarashi K., Sugino Y., Nishioka K.;

The complete nucleotide sequences of the cloned hepatitis B virus DNA; subtype adr and adw.";

DNA; subtype adr and adw.";

Nucleic Acids Res. 11:1747-1757(1983).
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PIR; C93460; NKVLA6.
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus
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InterPro; IPR002006; Hepatitis_core.
                                                                                       Hepatitis B virus (subtype adr4),
Hepatitis B virus (subtype adr), and
Hepatitis B virus (subtype adr), brain Indonesia/pIDW420)
Hepatitis B virus (subtype adr) of strain Indonesia/pIDW420)
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnaviru
NCBI_TaxID=10409, 106820, 10412;
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                                                                                                                                                                                                                                                  Core antigen.
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                                               SEQUENCE FROM N.A.
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177; Conserv
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172
185 AA;
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3; Mismatches
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MEDLINE=83168919;

PubMed=6300776;

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Best Local Sim
Matches 174;
                 CORA HPBVF STAN
P29178;
01-DEC-1992 (Rel. 2
01-DEC-1992 (Rel. 2
01-DEC-1992 (Rel. 2
                                                                                                                                     HPBVF
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EMBL; X01587; CAA25745.1; -
EMBL; D00331; -; NOT ANNOTATED CDS.
PIR; A93480; NKVLA5.
PIR; B93460; NKVLA4.
PIR; C28925; NKVLJ3.
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Okamoto H., Tsuda F., Sakugawa H., Sastros
Miyakawa Y., Mayumi M.;
"Typing hepatitis B virus by homology in r
comparison of surface antigen subtypes.";
J. Gen. Virol. 69:2575-2583(1988).
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Pfam; PF00906; Hepatitis_core; 1.
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Fujiyama A., Miyanohara A., Nozaki C.,
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                   (Rel. 24, Created)
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170
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                                                                                                             STANDARD;
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177
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92.6%;
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Pred. No. 2.9e-75;
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                                                                                          Query Match
Best Local S
Matches 176
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Best Local
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21-JUL-1986
01-AUG-1992
15-DEC-1998
                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                 Core protein; Repeat
REPEAT 164 1:
REPEAT 172 1:
                                                                                                                                                                                                                                                  InterPro; IPR002006; Hepatitis_core.
Pfam; PF00906; Hepatitis_core; 1.
                                                                                                                                                                                                                                                                                             Animal virus genetics,
PIR; A94409; NKVLA3.
                                                                                                                                                                                                                                                                                                              Valenzuela P., Quiroga M., Zaldi
(In) Field B.N., Jaenisch R., Fo.
Animal virus genetics, pp.57-70,
                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=10408;
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Viruses; Retroid viruses; Hepadn
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SEQUENCE
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Bhat R.A., Ulrich P.P., Vyas G.N.;
"Molecular characterization of a new variant
a persistently infected homosexual man.";
Hepatology 11:271-276(1990)
PIR; A37182; NKVLH3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORA_HPBV2
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REPEAT 174 181
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Viruses; Retroid viruses; Hepadnaviridae; O
NCBI_TaxID=31515;
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                           MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL 87
RRRSQSREPQC
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                                                                                                               Similarity
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74; Conservative
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172
185 AA;
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195 AA;
                                                                                            Conservative
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(Rel. 23, Last sec
(Rel. 37, Last ann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                    viruses, Hepadnaviridae, Orthohepadnavirus
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91.1%;
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                                                                                                                                                                         21304 MW;
                                                                                                         79.4%;
92.6%;
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R., Fox C.
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annotation update)
                                                                                        Score 911.5;
Pred. No. 1.2e
3; Mismatches
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                      J., Gray P., F. (eds.);
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                                                                                                           .2e-74;
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Best Local Sim
Matches 172;
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PIR; A28885; NKVLCP.
InterPro; IPR002006; Hepatitis core.
Pfam; PF00906; Hepatitis core; 1.
Core protein; Repeat.
REPEAT 162 169
REPEAT 170 177
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01-OCT-1989
01-OCT-1989
01-APR-1990
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CORA_HPBVL
                                                                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for contentitles requires a license agreement (See http://www.isb-sib.ch/aror send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                             virus isolated from a naturally in J. Gen. Virol. 69:1383-1389(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                       "The complete nucleotide sequence of the genome of a hepatitis virus isolated from a naturally infected chimpanzee.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vaudin M., Wolstenholme Harrison T.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis B virus (strain lsh / chimpanzee isolate).
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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MEDLINE=88258473; PubMed=2838576;
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                                                                                                                                                                                                                                          SEQUENCE
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                                                              CWGELMTLASWVGNNLEDP
                                                                                                           CWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQLLWFHISCLTFGRETV 147
                                                                                                                                            MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPNHTALRQAIL 60
SQSPASQC
                      SOSREPOC
                                              LEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRTPSPRRRRSQSPRRRR
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(Rel. 12,
(Rel. 14,
                                                                                                                                                                                                                                          183 AA;
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 183
                      215
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                                                                                                                                                                                                                                          20999 MW; 923DCB94A33FC0E8 CRC64;
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91.5%;
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                                                                                             ----ASREQVVNYVNTNMGLKIRQLLWFHISCLTFGRETV
                                                                                                                                                                                          Score 899.5;
Pred. No. 1.4e
5; Mismatches
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(See http://www.isb-sib.ch/announce/
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RESULT 13

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CORA HPBGS
CORA HPBGS
AC P03153
DT 21-UIL
DT 21-UIL
DT 01-UN
DE Core a
GN C.
OS Ground
OC Viruse
OX (N] I
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CORA_WHV1
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Query Match
Best Local S
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21-JUL-1986
21-JUL-1986
01-JAN-1990
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P03152;
21-JUL-1986 (Rel. (
21-JUL-1986 (Rel. (
15-DEC-1998 (Rel. )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=84267998; PubMed=6086950;
Seeger C., Ganem D., Varmus H.B.;
"Nucleotide sequence of an infectious molecularly cloned genome
ground squirrel hepatitis virus.";
J. Virol. 51:367-375(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR002006; Hepatitis_core. Pfam; PF00906; Hepatitis_core; 1. Core protein; Repeat.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=10406;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ground squirrel hepatitis virus (GSHV).
Viruses, Retroid viruses, Hepadnaviridae, Orthohepadnavirus
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        Woodchuck hepatitis virus 1 (WHV 1), Woodchuck hepatitis virus 7 (WHV 7), woodchuck hepatitis virus 59 (WHV 59), and Woodchuck hepatitis virus 8 (infectious clone) (WHV 8). Woodchuck hepatitis virus 8 (infectious clone) (WHV 8). Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus. NCBI_TaxID=10430, 10432, 10431, 10434;
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                                                                                                                                Core antigen
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204
217 AA;
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(Rel. 01, Last sequence up
(Rel. 13, Last annotation
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25189 MW; DF489467355EC11A CRC64;
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65.6%; Pr
Live 22;
                                                                                                                                              Created)
Last seq
Last anno
                                                                                                                                               sequence update) annotation updat
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Pred. No. 8.7e-61;
2; Mismatches 42
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EMBL; J04514; AAA46772.1; ...
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PIR; C32397; NKVLC4.
PIR; C32397; NKVLC4.
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SPECIES=Woodchuck hepatitis virus 1;

MEDLINE=82216969; PubMed=7086958;

Galibert F., Chen T.N., Mandart E.;

"Nucleotide sequence of a cloned woodchuck hepatitis virus comparison with the hepatitis B virus sequence.";

"Virol. 41:51-65(1982).
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SPECIES=Woodchuck hepatitis virus 7,

MEDLINE=88101359; PubMed=3336938;

Cohen J.I., Miller R.H., Rosenblum B.

Purcell R.H.;
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MEDLINE-89184524; PubMed=2928306;
Girones R., Cote P.J., Hornbuckle W.E., Tennant B.C., Gerin J.L.,
Purcell R.H., Miller R.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Sequence comparison of woodchuck hepatitis virus replicative forms shows conservation of the genome."; virology 162:12-20(1988).
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Kodama K., Ogasawara N., Yoshikawa H., Murakami S.;
"Nucleotide sequence of a cloned woodchuck hepatitis
evolutional relationship between hepadnaviruses.";
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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ALIGNMENTS

A;MOlecule type: DNA	A;Accession: S47405	A; Reference number: \$47404	A; Description: Molecular cloning and sequencing of two complete genomes of polish isoli	submitted to the EMBL Data Library, August 1994	R; Plucienniczak, A.	C; Accession: S47405; S53191; S53209; S53234; S53264; S53249; S53262; S53277; Anaria	C;Date: 18-Dec-1981 #sequence revision 08-Nov-1996 #text change 16-Jul-1999	cheri/83	A: Variety: subtype ayw4, isolate hb321: isolate patient Ferrague: 103, isolate patient	C;Species: hepatitis B virus, HBV	N; Contains: core antigen; e antigen	N;Alternate names: HBe antigen precursor / HBc antigen; pre-C/C antigen	e antigen precursor / core antigen - hepatitis B virus (subtype ayw4, isolate hb321 an	NKVLAH	RESULT 1	

A.Molecule type: DNA
A,Rolecule type: DNA
A,Residues: 1-212 < APLUS
A,Cross-references: EMBL:Z35716; NID:g527435; PIDN:CAA84786.1; PID:g527437
A,Experimental source: subtype ayw4, isolate hb321
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.
submitted to the EMBL Data Library, March 1995
A;Reference number: S53112
A;Accession: S53191

genomes of polish isolate

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A;Molecule type: DNA
A;Residues: 1-212 <LAI>
A;Residues: 1-212 <LAI>
A;Cross-references: EMBL:X85283; NID:g736088; PIDN:CAA59593.1; PID:g736090
A;Experimental_source: isolate patient Ferracuti'83 A; Accession: S53209

A;Molecule type: DNA
A;Residues: 1-212 <LR2>
A;Cross-references: EMBL:X85290; NID:g736114; PIDN:CAA59609.1; PID:g736116
A;Experimental source: isolate patient Castag'83
A;Accession: S53234

A; Molecule type: DNA A; Residues: 1-212 <LA3>

A;Cross-references: EMBL:X85300; NID:g736150; PIDN:CAA59631.1; PID:g736152
A;Experimental source: isolate patient Sanna'84
A;Accession: S5364
A;Molecule type: DNA
A;Residues: 1-212 <LA4>
A;Cross-references: EMBL:X85313; NID:g736194; PIDN:CAA59659.1; PID:g736196
A;Experimental source: isolate patient Licheri-1'85
A;Accession: S53249
A;Molecule type: DNA
A;Residues: 1-212 <LA5>
A;Cross-references: EMBL:X85306; NID:g736172; PIDN:CAA59644.1; PID:g736174
A;Experimental source: isolate patient Flore-1'86

A;Molecule type: DNA A;Residues: 1-212 <LA6> A;Cross-references: EMBL:X85312; NID:g736191; PIDN:CAA59657.1; PID:g736193 A;Experimental source: isolate patient Licheri'83

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NyALTERINATE HAME ANALYSES. PRECULSOL / NBC ANLIGHING PRECULSOL / NBC ANLIGHNA PRECULSOL / NBC ANLIGHNA PRECULSOL / NBC ANLIGHNA PRECULSOL / NBC ANLIGHNA PRECULSOL / NBC ANLIGHNA PRECULSOL / NBC ANLIGHNA PROPERTY.

C;Species hepatitis B virus, HBV

C;Species hepatitis B virus, HBV

C;Accession: 553204; 553283

R;La1, M.E., Mazzoleni, A.P.; Porru, A.; Balestrieri, A.

submitted to the EMBL Data Library, March 1995

A;Accession: 553204

A;Accession: 553203

A;Accession: 553204

A;Accession
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A,Experimental source: patient Giordo-2'86
A;Note: due to a stop codon between the alternative initiators the e antigen R;Galibert, F.; Mandart, E.; Fitoussi, F.; Tiollais, P.; Charnay, P.
Nature 281, 646-650, 1979
A;Title: Nucleotide sequence of the hepatitis B virus genome (subtype ayw) in A;Reference number: A93214; MUID:81012091; PMID:399327
A,Accession: A03711
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A;Residues: 1-212 <GAL>
A;Cross-references: GB:JO2203; NID:g329640; PIDN:AAA45489.1;
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A; Residues: 30-212
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Best Local &
;Gene: C
;Superfamily; hepatitis B virus core antigen
;Keywords: alternative initiators; core protein
;Leywords: signal sequence #status predicted <S:
;1-29/Domain: signal sequence #status predicted <G:
;30-212/Product: core antigen #status predicted <EAG>
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Pred. No. 2e-82;
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e antigen precursor / core antigen - hepatitis B virus (isolate patient Castag-1'85 - N',Alternate names: HBc antigen; HBe antigen precursor / HBc antigen; pre-C/C antigen C.Species: hepatitis B virus, HBV
A,Variety: isolate patient Castag-1'85; isolate patient Perracuti-1'89
C;Date: 08-Jul-1995 #sequence_revision 03-Aug-1995 #text_change 26-Aug-1999
C;Accession: S53211, S53197
R;Lai, M.B.; Marzoleni, A.P.; Porru, A.; Balestrieri, A.
A,Reference number: S53112
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A;Residues: 30-212 <LAM>
A;Residues: 30-212 <LAM>
A;Cross-references: EMBL:X85284; NID:g736095; PIDN:CAA59596.1;
A;Experimental source: isolate patient Ferracuti-1'89
A;Note: due to a stop codon between the alternative initiations
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A;Experimental source: isolate patient Castag-1'85
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C, Genetics:
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                                                                                                                                                                                                                                                                                                                                                 PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP
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PSPRRRRSQSPRRRRSQSREPQC
                                                                                                  LWEHISCLTEGRETVIEYLVSEGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT 192
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Pred. No. 3e-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 1037.5; DB Pred. No. 3.6e-82;
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G;Species: hepatitis B virus, HBV
A;Variety: subtype ayw, patient CI
C;Date: 20-Reb-1995 #sequence_revision 20-Reb-1995 #text_change 2
C;Accession: S20750
R;Lai, M.E.; Mazzoleni, A.P.; Balestrieri, A.; Melis, A.; Porru,
submitted to the EMBL Data Library, March 1992
A;Description: Sequence analysis of HBV genomes isolated from pat
A;Reference number: S20745
                                                                                                                                                                           e antigen precursor / core antigen - hep:
N;Alternate names: HBe antigen precursor
N;Contains: core antigen; e antigen
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A;Residues: 30-212 <LAI>
A;Residues: 30-212 <LAI>
A;Cross-references: EMBL:X85289; NID:g736110; PIDN:CAA59607.1;
A;Experimental source: isolate patient Castaa-2'87
A;Note: due to a stop codon mutation between the alternative in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: EMBL:X72702; NID:g288927; PIDN:CAA51257.1; A;Experimental source: subtype ayw, patient C1000 R;Lai, M.B.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A. submitted to the EMBL Data Library, March 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R:Preisler-Adams, S.; Schlayer, M.J.; Peters, T.; Hettler, F.; Gerok, W.; Rasenz submitted to the EMBL Data Library, March 1993
A;Description: Identification and sequence analysis of hepatitis B virus DNA in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e antigen precursor / core antigen - hepatitis B virus (subtype ayw, patients C1000 N,Alternate names: HBe antigen precursor / HBc antigen; pre-C/C antigen N;Contains: core antigen; e antigen C;Species: hepatitis B virus, HBV
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 20-Oct-2000 C;Accession; S32204; S53207
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A; Residues: 1-212 < PRE>
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                                                                                                                                                                                                                                                                                                                                                                                   LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT
                                                                                                                                                                                                                                                                                                                                                                                                            LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                            EHCSPHHTÄLRQAILCWGELMTLATWVGVNLEDP----ASRDLVVSYVNTNMGLKFRQL
                                                                                                                                                                                                                                                                                                               PSPRRRRSQSPRRRRSQSRESQC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDAGRDLVVSYVNTNMGLKFRQL 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFFSVRDLLDTASALYREALESP 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYRDALESP 74
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Pred. No. 3.6;
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C;Superfamily: hepatitis B virus core antigen C;Keywords: alternative initiators; core protein F;1-29/Domain: signal sequence #status predicted <8IG>F;30-212/Product: core antigen #status predicted <CAG>F;30-178/Product: eantigen #status predicted <EAG>F;30-178/Product: eantigen #status predicte
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C; Date: 08-Jul-1995 #sequence_revision
C; Accession: S53216
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S53216
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A;Experimental source: isolate patient Castag'3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              submitted to the EMBL Data A; Reference number: $53112 A; Accession: $53216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-212 < LAI>
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Alternate names: HBe antigen precursor / HBc
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LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT 192
                                                                                                                                                                                                                                    PTVQASKLCLGWLWGMDIDPYXBFGATVBLLSFLPSDFFPSVRDLLDTASALYRBALESP
                                                                EHCSPHHTALRQAILCWGBLMTLATWVGVNLEDP----ASRDLVVSYVNTNMGLKFRQL
                                                                                                            EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQL 132
                                                                                                                                                                                              PTVQASKLCLGWLWGWDIDPYKEFGATVELLSFLPSDFFPSVRDLLDNASALYREALESP
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Pred. No. 3.6e-82;
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e antigen precursor / core antigen - hepatitis B virus N,Alternate names: HBe antigen precursor / HBc antigen N;Contains: core antigen; e antigen C;Species: hepatitis B virus, HBV A,Variety: isolate patient Chighine-2'86 C;Date: 08-UL-1995 #sequence_revision 03-Aug-1995 #t6 C;Accession: 853225 R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, submitted to the EMBL Data Library, March 1995 A,Reference number: 853112 A,Reference number: 853112
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C/Superfamily: hepatitis B virus core antigen
C/Superfamily: hepatitis B virus core protein
C/Keywords: alternative initiators; core protein
F/1-29/Domain: signal sequence #status predicted <SIG>
F/30-212/Product: core antigen #status predicted <CAG>
F/30-178/Product: e antigen #status predicted <EAG>
F/30-178/Product: e antigen #status predicted #link EAG #status predicted
F/179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted
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A;Residues: 1-212 <LAI>
A;Crosa-references: EMBL:X85315; NID:g736205; PIDN:CAA59664.1; PID:g736207
A;Experimental source: isolate patient Licheri-3'90
C;Genetics:
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       A;Molecule type: DNA
A;Residues: 1-212 <LAI>
A;Residues: 1-212 <LAI>
A;Cross-references: EMBL:X85296; NID:g736137; PIDN:CAA59622.1; PID:g736139
A;Experimental source: isolate patient Chighine-2'86
A;Experimental source:
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Alternate names. HBe antigen precursor / HBc antigen; pre-C/C antigen
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Pred. No. 5.4e-82;
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PSPRRRRSQSPRRRRSQSREPQC
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A;Gene: C
;Superfamily: hepatitis B virus core antigen
C;Superfamily: hepatitis B virus core protein
C;Keywords: alternative initiators; core protein
C;Keywords: alternative initiators; core protein
E;1-29/Domain: signal sequence #status predicted <SIG>
E;30-212/Product: core antigen #status predicted <CAG>
F;30-178/Product: e antigen #status predicted <EAG>
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted
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A;Accession: $53274
A;Molecular
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                                                                                                                                                                                                                                                                                                                                                                                          C;Keywords: alternative initiators; core protein F;1-29/Domain: signal sequence #status predicted <SIG>F;30-212/Product: core antigen #status predicted <CAG>F;30-178/Product: core antigen #status predicted <EAG>F;30-178/Product: e antigen #status predicted <EAG>F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted
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A; Residues: 1-212 <LAI>
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195;
                           LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT 192
                                                                                                                         EHCSPHHTALROAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQL 132
                                                                                                                                                                                             PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP
                                                                                                                                                                                                                      PTVQASKLCLGWLWGWDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP 72
                                                                                                  EHCSPHHTALRQAILCWGELMTLATWVGVNLEDP----
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96.6%;
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Pred. No. 6.6e-82;
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                                                                                                                                                                                                                                                                                                Score 1034.5; DB 2;
Pred. No. 6.6e-82;
0; Mismatches 2;
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C)Superfamily: hepatitis B virus core antigen C;Keywords: alternative initiators; core protein F;1-29/Domain: signal sequence #status predicted F;30-212/Product: core antigen #status predicted
                                                                                                                                                                                                                                                                                                                         e antigen precursor / core antigen - hepatitis B virus (isolate patient Flore'85)
N;Alternate names: HBe antigen precursor / HBc antigen; pre-C/C antigen
N;Contains: core antigen; antigen
C;Species: hepatitis B virus, HBV
A;Variety: isolate B virus, HBV
A;Variety: isolate patient Flore'85
C;Date: 08-Vul-1955 #sequence_revision 03-Aug-1995 #text_change 26-Aug-1999
                                                                                                                            A;Molecule type: DNA
A;Residues: 1-212 <LAI>
A;Cross-references: EMBL:X85307; NID:g736175; PIDN:CAA59646.1;
A;Experimental source: isolate patient Flore'85
                                                                                                                                                                                                                                                                     C;Accession: S53251
R;Lai, M.B.; Mazzoleni, A.P.; Porru,
submitted to the EMBL Data Library,
                                                                                                                                                                                                                               A; Reference number: $53112
A; Accession: $53251
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C;Keywords: alternative initiators; core protein
F;1-29,Domain: signal sequence #status predicted <SIG>
F;30-212/Product: core antigen #status predicted <CAG>
F;30-178/Product: e antigen #status predicted <EAG>
F;30-179/Product: e antigen #status predicted <EAG>
F;10-179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted
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A;Experimental source: isolate patient Vittorina'92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-212 <LAI>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Variety: isolate patient Vittorina'92
C;Dariety: 08-Jul-1995 #sequence_revision 03-Aug-1995 #text_change 26-Aug-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e antigen precursor / core antigen - hepatitis B virus (isolate patient Vittorina'92)
N;Alternate names: HBe antigen precursor / HBc antigen; pre-C/C antigen
N;Contains: core antigen; e antigen
C;Species: hepatitis B virus, HBV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PSPRRRRSQSPRRRRSQSRESQC
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Pred. No. 6.66
                                                                                                                                                                                                                                                                       , A.; Balestrieri, A. March 1995
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       ^CAG
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C.Keywords: alternative initiators; core protein
F;1-29/Domain: signal sequence #status predicted <SIG>
F;30-212/Product: core antigen #status predicted <CAG>
F;30-178/Product: e antigen #status predicted <EAG>
F;30-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted
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A; Residues: 1-212 < LAI>
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C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change
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submitted to the RME Data Library, March 1992
A;Description: Sequence analysis of HBV genomes isolated
A;Reference number: S20745
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N;Contains: core antigen; e antigen
C;Species: hepatitis B virus, HBV
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                                                                                                                                                                                                                                                     Matches 195;
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                                                                        LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRGRSPRRFT
                          PSPRRRRSQSPRRRRSQSREPQC 215
                                                                                                                 EHCSPHITALROATLCWGELWTLATWVGVNLEDP----ASRDLVVSYVNTNWGLKFRQL
                                                                                                                                          EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNWGLKFRQL 132
                                                                                                                                                                                                  PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYRBALBSP
PSPRRRRSQSPRRRRSQSRESQC
                                                              LWFHISCLIFGRETVI
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96.1%;
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                                                                                                                                                                                                                                              Score 1031.5;
Pred. No. 1.2e
1; Mismatches
212
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Pred. No. 1.2e.
1; Mismatches
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                                                                                                                                                                                                                                                                            DB 2;
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RESULT

13

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e antigen precursor / core antigen - hepatitis B virus (isolate patient Garrucciu'90)
N,Alternate names: HBe antigen precursor / HBc antigen; pre-C/C antigen
N;Contains: core antigen; e antigen
C;Species: hepatitis B virus, HBV
A;Variety: isolate patient Garrucciu'90
C;Date: 08-Jul-1995 #sequence_revision 03-Aug-1995 #text_change 26-Aug-1999
C;Accession: S53159
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.
submitted to the EMBL Data Library, March 1995
A;Accession: S53112
A;Accession: S53112
A,Accession: S53112
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A;Rolecule type: DNA
A;Residues: 1-212 <LAI>
A;Cross-references: EMBL:X85272; NID:g736044; PIDN:CAA59563.1; PID:g736046
A;Cross-references: isolate patient Garrucciu'90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Gene: C
C;Superfamily: hepatitis B virus core antigen
C;Superfamily: hepatitis B virus core protein
C;Keywords: alternative initiators; core protein
E;1-29/Domain: signal sequence #status predicted <SIG>
F;30-212/Product: core antigen #status predicted <CAG>
F;30-178/Product: e antigen #status predicted <CAG>
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted
                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-212 <LAI>
A; Cross-references: EMBL: X85286; NID:g736101; PIDN:CAA59600.1; PID:g736103
A; Cross-referental source: isolate patient Ferracuti-3'91
C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e antigen precursor / core antigen - hepatitis B virus (isolate N;Alternate names: HBe antigen precursor / HBc antigen; pre-C/C N;Contains: core antigen; e antigen C;Species: hepatitis B virus, HBV A;Variety: isolate patient Ferracuti-3/91 A;Variety: 180-195 #sequence_revision 03-Aug-1995 #text_change C;Date: 08-041-1995 #sequence_revision 03-Aug-1995 #text_change
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R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri,
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                                                                                                 A;Gene: C
C;Superfamily: hepatitis B virus core antigen
C;Superfamily: hepatitis B virus core protein
C;Keywords: alternative initiators; core protein
C;Keywords: signal sequence #status predicted <SIG>F;1-29/Domain: signal sequence #status predicted <CAG>F;30-212/Product: core antigen #status predicted <EAG>F;30-178/Product: e antigen #status predicted <EAG>F;30-178/Product <EAG>F;30-178/Product <EAG>F;30-178/Product <EAG>F;30-178/Product <EAG>F;30-178/Product <EAG>F;30-178/Product <EAG>F;30-178/Product <EAG>F
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Best Local S
Matches 194
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Pred. No. 1.5e
3; Mismatches
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Score 1030.5;
                                                                    propeptide #link EAG #status predicted
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80
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e antigen precursor / core antigen - hepatitis B virus (isolate N;Alternate names: HBe antigen precursor / HBc antigen; pre-C/C N;Contains: core antigen; e antigen C;Species: hepatitis B virus, HBV A;Variety: isolate patient Castaa'83 C;Date: 08-041-1955 #sequence_revision 03-Aug-1995 #text_change C;Date: 08-041-1955 #sequence_revision 03-Aug-1995 #text_change
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submitted to the EMBL Data Library, March
A;Reference number: S53112
A;Accession: S53202
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C;Superfamily: hepatitis B virus core antigen
C;Superfamily: hepatitis B virus core protein
C;Keywords: alternative initiators; core protein
E;1-29/Domain: signal sequence #status predicted -
E;30-212/Product: core antigen #status predicted -
E;30-178/Product: e antigen #status predicted -
E;179-212/Domain: carboxyl-terminal propeptide #1
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Matches
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                                                           EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQL
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                                                                                                                                                                                                          PTVQASKLCLGWLWGWDIDPYKEFGATVELLSFLPSDFPPSVRDLLDTASALYREALESP
                                                                                                                           EHCSPHHTALROAILCWGELMTLATWVGVNLEDP---
                                PSPRRRRSQSPRRRRSQSREPQC
PSPRRRRSQSPRRRRSQSRESQC
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96.1%;
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1; Mismatches 2
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                                                                                                                                                                                                                                                       Score 1030.5; DB
Pred. No. 1.5e-81;
1; Mismatches 2
     212
                                 215
                                                                                                                                                                                                                                                                                                                       predicted <EAG>
propeptide #link EAG #status predicted
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March 1995
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                                                                                                                                -ASRDLVVSYVNTNMGLKFRQI
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Search completed: February Job time : 15.2334 secs

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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      Sequence 1, Appli
Sequence 2, Appli
Sequence 3, Appli
Sequence 3, Appli
Sequence 31, Appli
Sequence 6, Appli
Sequence 17, Appli
Sequence 17, Appli
Sequence 17, Appli
Sequence 19, Appli
Sequence 19, Appli
Sequence 674, Appli
Sequence 678, Appli
Sequence 679, Appli
Sequence 678, Appli
                                                                                                                                                                                                                                                                                                                                                                       RESULT 1

US-10-274-616-1

Sequence 1, Application US/10274616

GENERAL INFORMATION:

APPLICANT: Birkett, Ashley J.

TITLE OF INVENTION: INFLUENZA IMMUNOGEN AND VACE FILE REFERENCE: ICC 127.0 4564/88545

CURRENT APPLICATION NUMBER: US/10/274,616

CURRENT FILING DATE: 2002-10-21

PRIOR APPLICATION NUMBER: 09/930,915

PRIOR FILING DATE: 2001-08-15

PRIOR FILING DATE: 2001-08-15

PRIOR FILING DATE: 2002-02-21

INUMBER OF SEG ID NOS: 87

SOFTWARE: Patentin version 3.1

SEQ ID NO 1

LENGTH: 183

TYPE: PRT

ORGANISM: Hepatitis B virus
Sequence 4, Application US/10274616
GENERAL INFORMATION:
APPLICANT: Birkett, Ashley J
TITLE OF INVENTION: INFLUENCA IMMUNOGEN AND VACCINE
FILE REFERENCE: ICC 127.0 4564/88545
                                                              RESULT 2
US-10-274-616-4
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Best Local !
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96.8%;
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US-10-226-007-713
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US-10-226-007-608
US-10-226-007-688
US-10-226-007-723
US-10-226-007-657
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Pred. No. 7.4e-62;
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Result No.

Score

Match

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Description

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SUMMARIES

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US-10-274-616-1
US-10-274-616-3
US-10-274-616-3
US-10-274-616-3
US-10-274-616-3
US-10-274-616-6
PCT-US02-17536-17
PCT-US02-17536-18
US-10-274-616-5
US-09-863-054-19
US-10-226-007-646
US-10-226-007-646
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Minimum DB Maximum DB

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Database

Pending Patents AA New:*

1: /cgn2_6/ptodata/1/paa/PCT_NEW_COMB.pep:*

2: /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*

3: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

4: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

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6: /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*

7: /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Total number of hits satisfying chosen parameters:

404799 seqs, 80919614 residues

Scoring table:

BLOSUM62 Gapop 10.0 ,

Gapext 0.5

Title: Perfect score:

US-09-890-752A-2 1148 1 MPLSSIFSRIGDPTVQASKL......RRRRSQSPRRRRSQSREPQC

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protein search, using

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mode1

Copyright

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Ltd

February 3,

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US-10-274-616-2

Sequence 2, Application US/10274616

GENERAL INFORMATION:

APPLICANT: Birkett, Ashley J.

APPLICANT: Birkett, Ashley J.

FILE REFERENCE: ICC 127.0 4564/88545

CURRENT APPLICATION NUMBER: US/10/274,616

CURRENT FILING DATE: 2002-10-21

PRIOR APPLICATION NUMBER: 09/930,915

PRIOR APPLICATION NUMBER: 09/930,915

PRIOR APPLICATION NUMBER: 10/080,299

PRIOR FILING DATE: 2002-02-21

NUMBER OF SEQ ID NOS: 87

NUMBER OF SEQ ID NOS: 87
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; LENGTH: 183

; TYPE: PRT

; ORGANISM: Hepatitis B virus

US-10-274-616-4
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CURRENT FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: 09/930,915
PRIOR FILING DATE: 2001-08-15
PRIOR APPLICATION NUMBER: 10/080,299
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 87
SOFTWARE: Patentin version 3.1
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Best Local Similarity 93.6%;
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Best Local Similarity 93.2%;
Matches 177; Conservative
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                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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                                                                                                           CWGELMTLATWVGNNLQDP----ASRDLVVNYVNTNMGLKIRQLLWFHISCLTFGRETV 115
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Pred. No. 1.8e-60;
5; Mismatches 2
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pred. No. 1.5e-59;
3; Mismatches 3;
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                                                                                                                 ; TYPE: PRT
; ORGANISM: Hepatitis B virus
US-10-226-007-31
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GENERAL INFORMATION:
APPLICANT: Birkett, Ashley J.
TITLE OF INVENTION: INFLUENZA IMMUNOGEN AND VACCINE
FILE REFERENCE: ICC 127.0 4564/88545
CURRENT APPLICATION NUMBER: US/10/274,616
CURRENT FILING DATE: 2002-10-21
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best Local Similarity 92.6%;
Matches 176; Conservarian
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                                                                                                                                                                     SOFTWARE: PatentIn version 3.1
SEQ ID NO 31
LENGTH: 183
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                                   Query Match
Best Local Similarity
Matches 170; Conserv
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PRIOR FILING DATE: 2001-08-15
PRIOR APPLICATION NUMBER: 10/080,299
PRIOR APPLICATION NUMBER: 10/080,299
                                                                                                                                                                                                                         TITLE OF INVENTION: Compositions and Therapeutic Methods for Viral Infection FILE REFERENCE: 5005.01

CURRENT APPLICATION NUMBER: US/10/226,007

CURRENT FILING DATE: 2002-11-15

PRIOR APPLICATION NUMBER: US 60/313,883

PRIOR FILING DATE: 2001-08-21

NUMBER OF SEQ ID NOS: 1673
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                                                                                                                                                                                                                                                                                                                                         APPLICANT: Zavitz, Kenton
APPLICANT: Hobden, Adrian
TITLE OF INVENTION: Composi
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Myriad Genetics, APPLICANT: Morham, Scott
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ORGANISM: Hepatitis B virus
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                                       Conservative
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 Mismatches

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Pred. No. 5.8e-59;
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                                         Score 901.5; DB 6
Pred. No. 3.1e-58;
8; Mismatches 5
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; ORGANISM: Spermophilus variegatus
US-10-274-616-6
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US-10-274-616-6
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APPLICANT: Holoshitz, Joseph
APPLICANT: Holoshitz, Joseph
APPLICANT: Ling, Song
TITLE OF INVENTION: Methods and Compositions for the Treatment of Diseases Associated
TITLE OF INVENTION: Signal Transduction Aberrations
FILE REFERENCE: UM-07135
CURRENT APPLICATION NUMBER: PCT/USO2/17536
CURRENT APPLICATION NUMBER: PCT/USO2/17536
PRIOR APPLICATION NUMBER: 60/295,691
PRIOR APPLICATION NUMBER: 60/295,691
PRIOR PILING DATE: 2001-06-04
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17, Application PC/TUS0217536 GENERAL INFORMATION:
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Best Local :
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SEQ ID NO 6
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TITLE OF INVENTION: TRIFUTANA IMMUNOGEN AND VACCINE
FILE REPERENCE: ICC 127.0 4564/88545
CURRENT APPLICATION NUMBER: US/10/274,616
CURRENT FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: 09/930,915
PRIOR FILING DATE: 2001-08-15
PRIOR FILING DATE: 2001-08-25
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 87
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RRG-----RSPRRRTPSPRRRRSQSPRRRRSQSREPQC 215
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                          Sequence 5, Application US/10274616
GENERAL INFORMATION:
APPLICANT: Birkett, Ashley J.
TITLE OF INVENTION: INFLUENZA IMMUNOGEN AND VACCINE
FILE REFERENCE: ICC 127.0 4564/88545
CURRENT APPLICATION NUMBER: US/10/274,616
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RESULT 9
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; OTHER INFORMATION: Synthetic PCT-US02-17536-17
                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: PCT/US02/17536
CURRENT FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/295,691
PRIOR FILING DATE: 2001-06-04
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin version 3.1
SEQ ID NO 18
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                                                                                                                                                                                                                                            Matches 141;
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LENGTH: 169
TYPE: PRT
ORGANISM: Artificial Sequence
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APPLICANT: Ling, Song
TITLE OF INVENTION: Methods
TITLE OF INVENTION: Signal
FILE REFERENCE: UM-07135
                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 16
                                                                                                                                                                                                                                                                                                                   FEATURE:
OTHER INFORMATION: Synthetic
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                                                                                                                                                                                                                                                          Local Similarity
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                                       FHISCLTEGRETVIEYLVSFGVWIRTPDAYRDDNAPILSTLD 176
                                                                                                                          CWGELMTLATWVGVNLED-PEFRGD-----ASRDLVVSYVNTNMGLKFRQLLW 134
                                                                                                                                                                   MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDNASALYREALESPEHCSPHHTALRQAIL
                                                                                                     CWGELMTLATWVGGNLEDHKDILEDERAAVDTYCVDPISRDLVVSYVNTNMGLKFRQLLW
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                                                                                                                                                                                                                                     62.9%; Score 722.5; DB 1; 87.0%; Pred. No. 2.6e-45; ative 1; Mismatches 7;
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87.7%;
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Pred. No. 2.6e-45
1; Mismatches (
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US-09-863-054-21
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GENERAL INFORMATION:
APPLICANT: Chisari, Francis V.
TITLE OF INVENTION: Peptides for Inducing Cytotoxic T
Lymphocyte Responses to Hepatitis B Virus
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PRIOR APPLICATION NUMBER: 09/930,915
PRIOR FILING DATE: 2001-08-15
PRIOR APPLICATION NUMBER: 10/080,299
PRIOR PILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 87
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TYPE: PRT
ORGANISM: Marmota monax
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              116 QEFLVSFGVWIRTPAPYRPPNAPILSTLPEHTVIRRRGGARASRSPRRRTPSPRRRRSQS 175
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mes 124; Conserv
                                                                                                                    PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/935,898
FILING DATE: 26-AUG-1992
APPLICATION NUMBER: US 08/024,120
FILING DATE: 26-FEB-1993
APPLICATION NUMBER: US 08/396,283
FILING DATE: 27-FEB-1995
APPLICATION NUMBER: US 08/463,486
FILING DATE: 05-JUN-1995
NAME: Lockyer, Jean M.
REGISTRATION NUMBER: 44,879
REFERENCE/DOCKET NUMBER: 014740-000421US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTMARE: PATEURIT RELEASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/863,054
FILING DATE: 21-Msy-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CWDELTKLIAWMSSNITSEQV-----RTIIVNHVNDTWGLKVRQSLWFHLSCLTFGQHTV 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQLLWFHISCLTFGRETV 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRRRRSQ 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Townsend and Townsend and Crew LLF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: San Francisco
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TOPOLOGY: linear

HOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 21:
US-09-863-054-21
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US-09-863-054-19
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US-10-218-140-1758
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CURRENT APPLICATION NUMBER: US/10/218,140
CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 09/540,763
PRIOR FILING DATE: 2000-03-30
PRIOR FILING DATE: 2000-03-30
PRIOR APPLICATION NUMBER: 60/127,728
PRIOR FILING DATE: 1999-04-05
PRIOR APPLICATION NUMBER: 60/127,636
PRIOR APPLICATION NUMBER: 60/127,636
PRIOR PILING DATE: 1999-04-02
PRIOR FILING DATE: 1999-04-02
PRIOR FILING DATE: 1999-04-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/127,607
PRIOR FILING DATE: 1999-03-31
NUMBER OF SEQ ID NOS: 6322
SOFTWARE: CUTANALOT Version 1.0
SEQ ID NO 1758
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                       Sequence 19, Application US/09863054
GENERAL INFORMATION:
GENERAL TROCKMATION: Francis V.
TITLE OP INVENTION: Peptides for Inducing Cytotoxic T
Lymphocyte Responses to Hepatitis B Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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APPLICANT: Shinkets, Richard A.
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND POLYPEPTIDES
TITLE OF INVENTION: ENCODED THEREBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                        222 KPRRSPTLRYRRSRSRSP----RRRSRSPKRRSPSPKREKHRSKSPRRHRSRSRD 272
                                                                                                                                                                                                                                                                                                                                                                                                                                             / Match 9.9%;
Local Similarity 50.0%;
les 28; Conservative
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              ZIP: 94111-3834
COMPUTER READABLE FORM:
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                                                                                                                                                                         NUMBER OF SEQUENCES:
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STRANDEDNESS: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH: 20 amino acids
                                                                                                                ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
MEDIUM TYPE: Floppy disk
                                                     COUNTRY: USA
                                                                           STATE: California
                                                                                           CITY: San Francisco
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                                  94111-3834
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 6;
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CURRENT APPLICATION NUMBER: US/10/226,007
CURRENT PILING DATE: 2002-11-15
PRIOR APPLICATION NUMBER: US 60/313,883
PRIOR FILING DATE: 2001-08-21
NUMBER OF SEQ ID NOS: 1673
SOFTWARE: Patentin version 3.1
SEQ ID NO 714
LENGTH: 20
TYPE: PRT
ORGANISM: Hepatitis B virus
US-10-226-007-714
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US-10-226-007-714
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MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 19:
US-09-863-054-19
                                                                                                            Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Morham, Scott
APPLICANT: Zavitz, Kenton
APPLICANT: Hobden, Adrian
TITLE OF INVENTION: Compositions and Therapeutic Methods for Viral Infection
FILE REFERENCE: 5005.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Myriad Genetics, Inc. APPLICANT: Morham, Scott
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                        149 EYLVSFGVWIRTPPAYRPPN 168
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                                                                                     Local Similarity
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REGISTRATION NUMBER: 44,879
REFERENCE/DOCKET NUMBER: 014740-000421US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/396,283
FILING DATE: 27-FEB-195
APPLICATION NUMBER: US 08/463,486
FILING DATE: 05-JUN-195
ATTORNEY/AGENT INFORMATION:
EYLVSFGVWIRTPPPYRPPN 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA;
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TYPE: amino acid
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FILING DATE: 26-AUG-1992
APPLICATION NUMBER: US 08/024,120
FILING DATE: 26-FEB-1993
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FILING DATE: 21-May-2001
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                                                                                     Conservative
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                                                                          Score 110; DB 6; Length 20; Pred. No. 0.085; 0; Mismatches 1; Indels
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Search completed: February Job time : 17.6045 secs

3, 2003, 09:35:00

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GENERAL INFORMATION:

APPLICANT: Myriad Genetics, Inc.

APPLICANT: Myriad Genetics, Inc.

APPLICANT: Myriad Genetics, Inc.

APPLICANT: Zavitz, Kenton

APPLICANT: Zavitz, Kenton

APPLICANT: Hobden, Adrian

TITLE OF INVENTION: Compositions and Therapeutic Methods for Viral Infection

CURRENT APPLICATION NUMBER: US/10/226,007

CURRENT APPLICATION NUMBER: US 60/313,883

PRIOR APPLICATION NUMBER: US 60/313,883

PRIOR APPLICATION TOMBER: US 60/313,883

SOFTWARE: Patentin version 3.1

SQFIWARE: Patentin version 3.1

SQFIWARE: Patentin version 3.1
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PRIOR FLING DATE: 2001-08-21

NUMBER OF SEQ ID NOS: 1673

SOFTWARE: Patentin version 3.1

SEQ ID NO 646

LENGTH: 20

TYPE: PRT
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US-10-226-007-679
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                                                                             Query Match 9.5%; Score 109; Best Local Similarity 95.0%; Pred. No. 0. Matches 19; Conservative 0; Mismatche
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Best Local Similarity 95.0%;
Matches 19; Conservative
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GENERAL INFORMATION:
APPLICANT: Myriad Genetics, Inc.
APPLICANT: Morham, Scott
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CURRENT FILING DATE: 2002-11-15
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TITLE OF INVENTION: Compositions and Therapeutic Methods for Viral Infection
FILE REFERENCE: 5005.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                         154 FGVWIRTPPAYRPPNAPILS 173
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1 FGVWIRTPPPYRPPNAPILS 20
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Pred. No. 0
                                                                             Mismatches
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                                                                                                                  DB 6; Length 20,
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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                 4569144 seqs, 644733110 residues
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1148
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Cgn2_6/ptodata/1/paa/USO6_COMB.pep:*
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(cgn2_6/ptodata/1/paa/USO86_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO83_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO83_COMB.pep:*
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(cgn2_6/ptodata/1/paa/USO86_COMB.pep:*
(cgn2_6/ptodata/1/paa/USO96_COMB.pep:*
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

ወጠቅጠኝት	Result No.
1148 100.0 215 1048.5 91.3 212 1048.5 91.3 212 1048.5 91.3 212 1048.5 91.3 212 1048.5 91.3 212 1040.5 90.6 212	Score
91.3 91.3 91.3 91.3	Query
212 212 212 212 212	Query Match Length DB ID
1 222 222	DB
22 US-09-890-752A-2 21 US-09-791-537-6815 22 US-09-848-616-114 26 US-10-243-739-54 26 US-10-244-065-54 1 PCT-US97-15500-3	ID
Sequence 2, Appli Sequence 6815, Ap Sequence 114, Appl Sequence 54, Appl Sequence 54, Appli Sequence 3, Appli	Description

Query Match 100.0%; Score 1148; DB 22; Length 215; Best Local Similarity 100.0%; Pred. No. 9.1e-110; Matches 215; Conservative 0; Mismatches 0; Indels 0;

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ALIGNMENTS

GENERAL INFORMATION: APPLICANT: Hildt, Eberhard APPLICANT: Hofschneider, Peter TITLE OF INVENTION: Particles for Gene Therapy FILE REFERENCE: 107070-120 (VOS-013) CURRENT APPLICATION NUMBER: US/09/890,752A CURRENT APPLICATION NUMBER: US/09/890,752A CURRENT FILING DATE: 2001-08-03 PRIOR FILING DATE: 2001-02-04 PRIOR FILING DATE: 2000-02-04 PRIOR APPLICATION NUMBER: DE 199 04 800.2 PRIOR FILING DATE: 1999-02-05 NUMBER OF SEQ ID NOS: 21 SOFTWARE: PATENTIAL Version 3.1 SEQ ID NO 2 LENGTH: 215 TYPE: PRT ORGANISM: Artificial Sequence
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APPLICANT: Belonomix, Inc.
APPLICANT: Debe, Derek
APPLICANT: Debe, Derek
APPLICANT: Debe, Derek
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES OF
TITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: 261/210
CURRENT APPLICATION NUMBER: US/09/791,537
CURRENT APPLICATION NUMBER: US/09/791,537
CURRENT FILING DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 153055
SOFTWARE: Patentin version 3.0
SEQ ID NO 6815
LENGTH: 212
TYPE: PRT
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Sebbel, Peter
APPLICANT: Sebbel, Peter
APPLICANT: Bachmann, Martin
APPLICANT: Bachmann, Martin
APPLICANT: Hechner, Franziska
TITLE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0180002
CURRENT APPLICATION UMBER: US/09/848,616
CURRENT ETLING DATE: 2001.05-05
NUMBER OF SEQ ID NOS: 186
SOFTWARE: Patentin Ver: 2.1
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US-09-791-537-6815
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Best Local Similarity 97.5
Matches 198; Conservative
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97.5%; Pred. No. 1.8e-99;
tive 0; Mismatches 0
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APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: In vivo Activation of Antigen Presenting Cells for Enhancement of
TITLE OF INVENTION: Inmune Responses Induced by Virus Like Particles
TITLE OF INVENTION. Inmune Responses Induced by Virus Like Particles
TITLE REFERENCE: 1700.021001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: PatentIn version 3.1
SEQ ID NOS 54
LENGTH: 212
TYPE: PRT
ORGANISM: Hepatitis B virus
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                  RESULT 5
US-10-244-065-54
; Sequence 54, A
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LENGTH: 212
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Best Local Similarity 97.5%;
Matches 198; Conservative
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GENERAL INFORMATION:
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TYPE: PRT
ORGANISM: Hepatitis B virus
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198; Conser
                      Application US/10244065
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nilarity 97.5%;
Conservative
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Pred. No. 1.8e-99
0; Mismatches
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Pred. No. 1.8e-99; 
0; Mismatches 0
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PRIOR APPLICATION NUMBER: 60/318,994
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOPTWARE: PatentIn version 3.1
SEQ ID NO 54
LENGTH: 212
Type: PRT
ORGANISM: Hepatitis B virus
US-10-244-065-54
                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 6
PCT-US97-15500-3
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Best Local Similarity 97.5%;
Matches 198; Conservative
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APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Pumpens, Paul
APPLICANT: Cielens, Indulis
APPLICANT: Renhofa, Regina
TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Particl
TITLE OF INVENTION: Method of Preparation and Use
PILE REFERENCE: 1700.0220001
                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows
SOPTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US97/15500
FILING DATE: 03-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/374,145
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CURRENT FILING DATE: 2002-09-16
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APPLICANT: Storni, 7
APPLICANT: Maurer, 1
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ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                    APPLICANT: The General Hospital Corporation TITLE OF INVENTION: INHIBITION OF VIRAL REPLICATION NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       193 PSPRRRRSQSPRRRRSQSREPQC 215
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                                                                                                                                                                                                          COUNTRY:
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         APPLICATION DATA:
                                                                                                                                                                                         RY: USA
02110-2804
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Tissot, Alain
Schwarz, Katrin
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; OTHER INFORMATION: Hepatitis B virus C antigen and e antigen ; OTHER INFORMATION: (HECAG/HBeAg) sequence US-09-705-547-11
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US-09-705-547-11
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                                                                                                                             Matches 197;
                                                                                                                                             Query Match
Best Local
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SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 212 amin
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Best Local Similarity 97.0%;
Matches 197; Conservative
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APPLICANT: Catharina Hultgren
IITLE OF INVENTION: WACCINGS CONTAINING RIBAVIRIN AND
IITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: TRIPEP.023AUS
CURRENT APPLICATION NUMBER: US/09/705,547
CURRENT FILING DATE: 2000-11-03
PRIOR APPLICATION NUMBER: 60/229,175
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                            LENGTH: 212
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REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 08-
TELECOMMUNICATION, INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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73 EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNNGLKFRQL 132
                                                   15
                                                                 13 PTVQASKLCLGWIWGMDIDPYKBFGATVELLSFLPSDFFPSVRDLLDTASALYRBALESP 72
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TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 617/542-5070
TELEPAX: 617/542-8906
                                       PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EHCSPHHTÄLROÄILCWGELMTLÄTWVGVNLEDP----ASRDLVVSYVNTNMGLKFROL
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                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                             FastSEQ for Windows Version 4.0
                                                                                                                           Conservative
                                                                                                                     90.6%; Score 1040.5; DB
97.0%; Pred. No. 1.2e-98;
tive 0; Mismatches 1
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                                                                                                                                                          DB 21; Length 212;
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APPLICANT: Blonomix, Inc.
APPLICANT: Blonomix, Inc.
APPLICANT: Debe, Derek
APPLICANT: Debe, Derek
APPLICANT: Debe, Derek
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES OF
TITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: 261/210
CURRENT APPLICATION NUMBER: US/09/791,537
CURRENT FILING DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 153055
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 92352
LENGTH: 212
TYPE: PAT
ORGANISM: Hepatitis B virus
US-09-791-537-92352
        Sequence 11. Application US/09929955

Sequence 11. Application US/09929955

GENERAL INFORMATION:
APPLICANT: Matti Sallberg

APPLICANT: Catharina Hultgren
TITLE OF INVENTION: VACCINES CONTAINING RIBAVIRIN AND
TITLE OF INVENTION: METHODS OF USE THEREOF
FILTE OF INVENTION: METHODS OF USE THEREOF
FILTE REFERENCE: TRIPEP.23AUS:
CURRENT APPLICATION NUMBER: US/09/929,955
CURRENT APPLICATION NUMBER: 09/705,547
PRIOR APPLICATION NUMBER: 09/705,547
PRIOR FILING DATE: 2000-108-15
PRIOR APPLICATION NUMBER: 60/229,175
PRIOR APPLICATION NUMBER: 60/229,175
PRIOR APPLICATION NUMBER: 60/225,767
PRIOR APPLICATION NUMBER: 60/225,767
PRIOR PILING DATE: 2000-08-17
NUMBER OF SEQ ID NOS: 49
SOFTWARE: FBSUSEQ for Windows Version 4.0
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US-09-791-537-92352
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US-09-929-955-11
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SOFTWARE: F
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                                                                                                                                                                                                                                                                                                                                                         212
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TITLE OF INVENTION: WETHODS OF USE THEREOF
TITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: TRIPPE, 23AUSC1
CURRENT APPLICATION UNMBER: US/10/104,966
CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: 09/705,547
PRIOR APPLICATION NUMBER: 09/705,547
PRIOR APPLICATION NUMBER: 09/29,175
PRIOR APPLICATION NUMBER: 60/29,175
PRIOR APPLICATION NUMBER: 60/29,175
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 15
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 212
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OTHER INFORMATION:
OTHER INFORMATION:
US-09-929-955-11
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; FEATURE:
; OTHER INFORMATION: Hepatitis B virus C antigen and
; OTHER INFORMATION: (HBCAG/HBeAg) sequence
US-10-104-966-11
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Best Local &
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GENERAL INFORMATION:
APPLICANT: Matti Sallberg
APPLICANT: Catharina Hultgren
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Best Local Similarity
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TYPE: PRT
ORGANISM: Artificial Sequence
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les 197; Conservative
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PSPRRRRSQSPRRRRSQSREPQC 215
                                                               LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT 192
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                                                                                                                         EHCSPHHTALROAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFROL 132
                                              197;
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                                                                                                                                                                                                                                90.6%; Score 1040.5; DB
97.0%; Pred. No. 1.2e-98;
tive 0; Mismatches
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Pred. No. 1.2e-98;
0; Mismatches 1
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US-09-791-537-125025
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OTHER INFORMATION: Hepatitis B virus C antigen and e
OTHER INFORMATION: (HBcAg/HBeAg) sequence
US-60-229-175-11
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                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 125025
LENGTH: 212
TYPE: PRT
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Best Local Similarity 96.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Bionomix, Inc.
APPLICANT: Debe, Derek
APPLICANT: Debe, Joseph
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES OF PROTEIN FAMILIES AND FAMILY MEMBE
TITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: 261/210
CURRENT APPLICATION NUMBER: US/09/791,537
CURRENT FILING DATE: 2001-02-22
RUMBER OF SEQ ID NOS: 153055
SOFTWARE: DEC ID NOS: 153055
SOFTWARE SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SECRETARY OF SE
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CURRENT FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 212
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GENERAL INFORMATION:
APPLICANT: Matti Sallberg
APPLICANT: Matti Sallberg
APPLICANT: Catherina Hultgren
TITLE OF INVENTION: VACCINES CONTAINING RIBAVIRIN
TITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: TRIPEP.023PR
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Best Local Similarity
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                                                                                                                                    Score 1038.5;
Pred. No. 1.9e.
1; Mismatches
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                                                                                                                                                               5; DB 21; Length .9e-98;
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Sequence 15944, Application US/09791537
GENERAL INFORMATION:
APPLICANT: Bionomix, Inc.
APPLICANT: Bionomix, Inc.
APPLICANT: Dabe, Derek
APPLICANT: Danzer, Joseph
ITITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES OF ITITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: 261/210
CURRENT APPLICATION NUMBER: US/09/791,537
CURRENT FILING DATE: 2001-02-2
NUMBER OF SEQ ID MOS: 153055
NUMBER OF SEQ ID MOS: 153055
SEO IN NO 15045
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US-09-791-537-15944
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GENERAL INFORMATION:
APPLICANT: STUTYUER Lieven
APPLICANT: UAN GEYT Caroline
APPLICANT: DE GENDT Sija
TITLE OF INVENTION: New HBV Sequences
FILE REFERENCE: 2551-52
CURRENT FILING DATE: 2000-11-22
PRIOR APPLICATION NUMBER: EP99870252.6
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: US/09/18,095
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 1999-12-07
VNUMBER OF SEQ ID NOS: 169
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SEQ ID NO 12

; LENGTH: 212

; TYPE: PRT

; ORGANISM: Hepatitis B virus

US-09-718-095-12
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SEQ ID NO 15944
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Pred. No. 2.4e+98;
1; Mismatches 1; I
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Sequence 47294, Application US/09791537
; GENERAL INFORMATION:
APPLICANT: Bionomix, Inc.
APPLICANT: Bionomix, Inc.
APPLICANT: Danzer, Joseph
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES OF PROTEIN FAMILIES AND FAMILY MEMBE
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES OF PROTEIN FAMILY MEMBE
TITLE OF INVENTION: USE THEREOF
FILE REFERENCE: 261/210
CURRENT APPLICATION NUMBER: US/09/791,537
CURRENT APPLICATION NUMBER: US/09/791,537
CURRENT FILING DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 153055
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 47294
LENGTH: 212
TYPE: PRT
ORGANISM: Hepatitis B virus
US-09-791-537-47294
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; TYPE: PAT
; ORGANISM: Hepatitis B virus
US-09-791-537-15944
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Search completed: February 3, 2003, 09:34:08 Job time : 154.571 secs
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                                                                                                                                                                                                                                                                                                                  Query Match 90.4%; Score 1037.5; DB 21; Length 212; Best Local Similarity 96.6%; Pred. No. 2.4e-98; Matches 196; Conservative 1; Mismatches 1; Indels 5;
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                                                                PSPRRRRSQSPRRRRSQSRESQC
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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1: /cgn2_6/ptodata/2/pubpaa/USO8_NEW_PUB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/PCT NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/USO6_PUBCOMB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*

7: /cgn2_6/ptodata/2/pubpaa/PCTUS_PUBCOMB.pep:*

8: /cgn2_6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*

9: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

10: /cgn2_6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*

11: /cgn2_6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*

12: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

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15: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

16: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

17: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
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        February 3, 2003, 09:25:53 , Search time 7.86585 Seconds (without alignments) 551.547 Million cell updates/sec
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US-10-068-059-6
US-10-068-059-6
US-10-068-059-6
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Sequence 12, Appl
Sequence 12, Appl
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Query M. Best Los Matches 13 15 73 75 133	SSULT 1 Sequence 11. Patent No. 1. Patent No. 1. GENERAL INFA APPLICANT: APPLICANT: TITLE OF IT FILE REFERI CURRENT APPL PRIOR FILLT PRIOR APPLL PRIOR APPLL PRIOR FILLT NUMBER OF SE SOFTWARE: F SOFTWARE: F SOFTWARE: F STATURE INFO OTHER INFO	44444000000000000000000000000000000000
atch cal s 197 PTVC PTVC - PTVC EHCS EHCS EHCS - EHCS	He 11, Application usage and the second seco	114 1114 1112 1111 1111 1111 1111 1111
Similarity 90.6%; Score 97.0%; Pred. 97.0%;		00000000000000000000000000000000000000
90.6% 97.0% Vative Adductor Ad	ilon US/101049 ilberg na Hultgren vaccines CONT METHODS OF U BEP.23AUSC1 NUMBER: US/10 2002-03-22 MBER: 09/705, 0000-11-03 MBER: 60/229, 1000-08-29 11 Sequence ll Sequence Hepatitis B v (HBCAg/HBeAg)	20 20 20 20 20 20 20 20 20 20 20 20 20 2
YKE YKE YKE YKE SFG	/101 % CO 985 003 985 003 100/2 29/7 118	100 100 100 100 100 100 100 100 100 100
1040.5; DB 9; L NO. 2.4e-95; NO. 2.4e-95; Insmatches 1; In LLSFLPSDFFPSVRDLLD LLSFLPSDFFPSVRDLLD LLSFLPSDFFPSVRDLUD NLEDPEFFGDASRDLVVS NLEDPASRDLVVS NLEDPASRDLVVS PAYRPPNAPILSTLPETT PAYRPPNAPILSTLPETT	/10104966 gyren BS CONTAINING RIBAVIRIN AND BS OF USE THEREOF USC1 : US/10/104,966 -03-22 09/705,547 -03 60/229,175 -29 compare Version 4.0 since since since iis B virus C antigen and e ant (HBeAg) sequence	0 US-09-839-447A-107 US-09-839-447A-111 US-09-839-447A-104 US-09-466-035-84 US-09-839-447A-105 US-09-839-447A-105 US-09-839-447A-106 US-09-839-447A-106 US-09-839-447A-106 US-09-839-447A-108 US-09-839-447A-108 US-09-839-447A-108 US-09-839-447A-109 US-09-839-417A-109 US-09-839-417A-109 US-09-839-417A-109 US-09-839-417A-109 US-09-839-417A-109 US-09-839-417A-109 US-09-839-417A-109 US-09-839-417A-109 US-09-839-417A-109 US-09-846-015-86 US-09-846-761-35898 US-10-151-736-4
Length 212; Indels 5; Gaps LLLDTASALYREALESP 72 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 72 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 72 LLDTASALYREALESP 74 LLDTASALYREALESP 72 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 74 LLDTASALYREALESP 72 LLDTASALYREALESP 74 LL	itigen	Sequence 107, App Sequence 81, App Sequence 84, App1 Sequence 101, App Sequence 105, App Sequence 106, App Sequence 106, App Sequence 99, App1 Sequence 98, App1 Sequence 110, App Sequence 110, App Sequence 109, App Sequence 109, App Sequence 107, App Sequence 107, App Sequence 107, App Sequence 108, App Sequence 107, App Sequence 108, App Sequence 108, App Sequence 107, App Sequence 108, App Sequence 109, App Sequence 107, App Sequence 108, App Sequence 108, App Sequence 109, App Sequence 13, App Sequence 37015, A Sequence 37015, A Sequence 4, Appli
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PSPRRRRSQSPRRRRSQSREPQC 215

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CURRENT APPLICATION NUMBER: US/09/929,955
CURRENT FILING DATE: 2001-08-15
PRIOR APPLICATION NUMBER: 09/705,547
PRIOR FILING DATE: 2000-11-03
PRIOR APPLICATION NUMBER: 60/229,175
PRIOR ETLING DATE: 2000-08-29
PRIOR APPLICATION NUMBER: 60/225,767
PRIOR PILING DATE: 2000-08-17
PRIOR PILING DATE: 2000-08-17
NUMBER: 09 SEQ ID NOS: 49
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 212
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US-10-209-264-4
(Sequence 4, Application US/10209264
(Publication No. US20030003111A1
(Publication NORMATION: Chong Jin
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Patent No. US200
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Best Local Similarity
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TITLE OF INVENTION: VACCINES CONTAINING RIBAVIRIN
TITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: TRIPEP. 23AUS2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Matti Sallberg APPLICANT: Catharina Hul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence FEATURE:
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CORRESPONDENCE ADDRESS:
ADDRESSE: Ladda & Parry
STREET: 26 West 61 Street
CITY: New York
STATE: New York
                                                                                                                                         Zhao, Yi
Chen, Wei Ning
TITLE OF INVENTION: A MUTANT HUMAN HEPATITIS B VIRAL STRAIN AND
                                                                                                                                                                                                        APPLICANT: Oon, Chong Jir
Lim, Gek Keow
                                                                                                     NUMBER OF SEQUENCES:
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Pred. No. 2.4e
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US-09-812-862-12
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APPLICATION NUMBER: PCT/SG98/00046

FILING DATE: 19-JAM-1998

ATTORNEY/AGENT INFORMATION:

NAME: Mass, Clifford J.

REGISTRATION NUMBER: 30,086

REFERENCE/DOCKET NUMBER: U-013109-7

TELECOMMUNICATION INFORMATION:

TELEPHONE: (212) 708-1800

INFORMATION FOR SEO ID NO: 4:

SEQUENCE CHARACTERISTICS:
LENGTH: 212 amino acids
TYPE: amino acid

TYPE: amino acid

TYPE: amino acid

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Patent No. US20020035081A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   130 LWFHISCLTFGRETVLEYLVSFGVWIRTPTAYRPPNAPILSTLPETTVVRRRGRSPRRRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        133 LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      190 PSPRERRSQSPRERRSQSRESQ 211
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            73 EHCSPHHTALRQAILCWGELMTLATWYGVNLEDPEFRGDASRDLVVSYVNTNWGLKFRQL 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNDER: US/10/209,264
PILING DATE: 31-Jul-2002
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10023
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                    Scaglioni, Pier Paolo
Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE DESCRIPTION: SEQ ID NO: 4:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EHCSPHHTALRQAILCWGELMNLATWVGSNLEDP----ASRELVVSYVNVNWGLKLRQI 129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
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                                                                                                                                                                             COUNTRY: USA
ZIP: 02110-2804
                                                                                                                                                                                                                                       STATE: MA
                                                                                                                                                                                                                                                                   CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      85.1%; Score 977.5; DB 9 91.1%; Pred. No. 3.9e-89; tive 5; Mismatches 8
                                          Version #1.30
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US-09-812-862-6
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/ SEQUENCE DESCRIPTION: SEQ ID NO: 12:
US-09-812-862-12
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Patent No. US20020035081A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 180; Conserv
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/09/812,862

PILING DATE: 20-Mar-2001

CLASSIFICATION: «Unknown»
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TELEX: 200154
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                            Scaglioni, Pier Paolo
Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/282001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/667,073
FILING DATE: «Unknown»
APPLICATION NUMBER: 08/492,489
FILING DATE: 20-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRTPSFRRRRSGSPRRRR 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKEROLLMFHISCLTFGRETV 147
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                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                       STATE: MA
                                                                                                                                                                                                                                      CITY: Boston
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TYPE: amino acid
TOPOLOGY: linear
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FILING DATE: 20-Mar-2001
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95.7%;
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Pred. No. 2.2
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GENERAL INFORMATION:

APPLICANT: Mizzen, Lee A.

APPLICANT: Mizzen, Liu

APPLICANT: Siegel, Marvin

ITILE OF INVENTION: HEBATITIS B VIRUS TREATMENT

FILE REFERENCE: 12071-017002

CURRENT APPLICATION NUMBER: US/10/068,059

CURRENT FILING DATE: 2002-06-04

PRIOR APPLICATION NUMBER: US 60/266,733

PRIOR FILING DATE: 2001-02-05

NUMBER OF SEQ ID NOS: 12

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 12

LENGTH: 724
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                                                                                                                                                    ; OTHER INFORMATION: Fusion protein US-10-068-059-12
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                                                                             Matches
                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 12, Application UPatent No. US20020155434A1
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                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                          TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 200154
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           176 SQSR 179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      148 IEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRTPSPRRRRSQSPRRRR 207
               28 MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 CWGELMTLATWVGVNLEDP-----ASRDLVVSYVNTNWGLKFRQLLWFHISCLTFGRETV 115
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mes 179; Conserv
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REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617,542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 6:
MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/492,489
FILING DATE: 20-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 397 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQLLWFHISCLTFGRETV 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/10068059
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                                                                            Conservative
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                                                                                           80.9%;
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97.3%;
                                                                      Score 928.5; DB 9;
Pred. No. 1.2e-83;
2; Mismatches 2;
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Pred. No. 1.8e-84;
0; Mismatches 0
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                                                                                                          DB 9;
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                                                                                                          Length 724;
                                                                          Indels
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                                                                    Gaps
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GENERAL INFORMATION:

APPLICANT: Mizzen, Lee A.

APPLICANT: Mizzen, Liu

APPLICANT: Hongwei, Liu

APPLICANT: Siegel, Marvin

TITLE OF INVENTION: HEPATITIS B VIRUS TREATMENT

FILE REFERENCE: 12071-017002

CURRENT APPLICATION NUMBER: US/10/068,059

CURRENT FILING DATE: 2002-06-04

PRIOR APPLICATION NUMBER: US 60/266,733

PRIOR FILING DATE: 2001-02-05

NUMBER OF SEQ ID NOS: 12

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 6

SEQ ID NO 6

TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Fusion protein

US-10-068-059-6
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US-10-068-059-6
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Sequence 6, Application US/10068059
Patent No. US20020155434A1
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Best Local Similarity 94.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches 179; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           176
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                                                                                                                                                                                                                                       RRSOSRESQC 205
                                                                                                                                                                                                                                                                                                                                                     CMGELMTLATWYGNNLEDP-----ASRDLVVNYVNTNMGLKFRQLLWFHISCLTFGRETV 135
                                                                                                                                                                                                                                                                 RRSOSREPQC 215
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; Pred. No. 1.3e-
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .3e-83;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 746;
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Sequence 2, Application US/10068059

patent No. US20020155434A1

general Impormation:

APPLICANT: Mizzen, Lee A.

APPLICANT: Mizzen, Liu

APPLICANT: Siegel, Marvin

ITILE OF INVENTION: HERATITIS B VIRUS TREATMENT

FILE REFERENCE: 12071-017002

CURRENT APPLICATION NUMBER: US/10/068,059

CURRENT FILING DATE: 2002-06-04

PRIOR APPLICATION NUMBER: US 60/266,733

PRIOR FILING DATE: 2001-02-05

NUMBER OF SEQ ID NOS: 12

SOPTMARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                             ; TYPE: PRT; ORGANISM: Hepatitis
US-10-068-059-2
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SOFTWARE: PatentIn version
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 2
LENGTH: 185
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Best Local Similarity
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Matches
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TYPE: PRT
ORGANISM: Hepatitis B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            148
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176
                           206 RRSOSREPOC 215
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nes 178; Conservative
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                                                        LEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRDRGRSPRRRTPSPRRRRSQSPRR
                                                                                                                                                                        | ISYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRR--RGRSPRRRTPSPRRRRSQSPRR 205
                                                                                                                                  CWGELMTLATWYGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQLLWFHISCLTFGRETV 147
 RRSQSRESQC
 185
                                                                                                                                                                                                                                                    80.4%;
                                                                                                                                                                                                                                                                                                              B virus
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94.7%;
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Pred. No. 3.4e-84;
                                                                                                                                                                                                                                    Score 922.5; DB 9;
Pred. No. 8.6e-84;
2; Mismatches 3;
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RESULT 10

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APPLICANT: Renner, Wolfgang A.

TITLE OF INVENTION: Compositions for Inducing Self-Specific Anti-IgE
TITLE OF INVENTION: Antibodies and Uses Thereof
FILE REFERENCE: 1700.0140001
CURRENT APPLICATION NUMBER: US/09/916,230
CURRENT FILING DATE: 2001-07-27
PRIOR APPLICATION NUMBER: US 60/221,841
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 35
SOCTWARE: PATENDER: US 60/221,841
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 35
SOCTWARE: PATENDER: US 60/221,841
CREANISM: Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-812-862-8
Sequence 8, Application US/09812862
Patent No. US20020035081A1
GENERAL INFORMATION:
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      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/812,862
PILING DATE: 20-Mar-2001
CLASSIFICATION - <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/667,073
APPLICATION NUMBER: 08/492,489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            176 RRSQSRESQC 185
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Local Similarity 93.7%;
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                                                                                                                                                              COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Wands, Jack R.
Scaglioni, Pier Paolo
Scaglioni, Pier Paolo
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CWGELMTLATWVGNNLEDP-----ASRDLVVNYVNTNMGLKIRQLLWFHISCLTFGRETV 115
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                                                                                                                                                                                                                                                                                                                  CITY: Boston
STATE: MA
FILING DATE: 20-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Fish & Richardson P.C
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Pred. No. 8.6e-84;
                                                                                                                                                                                                                                                                                                                                                        Street
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RESULT 12
US-09-812-862-4
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Patent No. US20020035081A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 176; Conserv
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INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
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                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/667,073
FILING DATE: <Unknown>
APPLICATION NUMBER: 08/492,489
FILING DATE: 20-JUN-1995
ATTORNEY/AGENT INFORMATION;
                                                                                                                                                                                                                                              ZIP: 02110-2804
COMPUTER READABLE FORM:
COMPUTER: Flappy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scaglioni, Pier Paolo
Melegari, Margherita
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/282001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IBYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRTPSPRRRRSQSPRRRR
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NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REPERENCE/DOCKET NUMBER: 00786/282001
                                                                                                                                                                                APPLICATION NUMBER: US/09/812,862
FILING DATE: 20-Mar-2001
CLASSIFICATION: -Unknown>
                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Boston
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96.2%;
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Pred. No. 2.4e-83;
1; Mismatches 1
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TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070

TELEFAX: 617/542-8906 TELEX: 200154

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LENGTH: 351 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-812-862-4
                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 12071-017002

CURRENT APPLICATION NUMBER: US/10/068,059

CURRENT FILING DATE: 2002-06-04

PRIOR APPLICATION NUMBER: US 60/266,733

PRIOR FILING DATE: 2001-02-05

NUMBER OF SEQ ID NOS: 12

SOFTWARE: PastSEQ for Windows Version 4.0

SEQ ID NO 10

LENGTH: 690
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Best Local Similarity 87.5%;
Matches 168; Conservative
                                                                                                                                                                                                                         Query Match
Best Local S
                                                                                                                                                                                                            Matches 148;
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APPLICANT: Siegel, Marvin
TITLE OF INVENTION: HEPATITIS B VIRUS TREATMENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Mizzen, Lee A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
                        4 148 IEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRR 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      179 QSPTŚNHSPTSC 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    204 RRRRSOSREPOC 215
116 LEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRR 151
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                                                                                                                                                                                                                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IBYLVSFGVWIRTPPAYRPPNAPILSTLPEHTVIRRGGARASRSPRRRTPSPRRRRSQNS 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CWGELMTLATWVGVNLEDP----ASRDLVVSYVNTNMGLKFRQLLWFHISCLTFGRETV
                                                                                          CWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQLLWFHISCLTFGRETV 147
                                                                      CWGELMTLATWVGNNLEDP----ASRDLVVNYVNTNMGLKFRQLLWFHISCLTFGRETV 115
                                                                                                                                         MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL 60
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ilarity 94.9%;
Conservative :
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Pred. No. 1.3e-77;
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Pred. No. 6.9e-69;
2; Mismatches 1
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Sequence 4, Application US/10068059

Patent No. US20020155434A1

; GENERAL INFORMATION:
APPLICANT: Mizzen, Lee A.
APPLICANT: Hongwei, Liu
APPLICANT: Hegel, Marvin
TITLE OF INVENTION: HEPATITIS B VIRUS TREATMENT
TITLE OF INVENTION HEPATITIS B VIRUS TREATMENT
CURRENT APPLICATION NUMBER: US/10/068,059
; CURRENT FILING DATE: 2002-06-04
PRIOR APPLICATION NUMBER: US 60/266,733
; PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 12
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 4
LENGTH: 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Mizzen, Lee A.
APPLICANT: Hongwei, Liu
APPLICANT: Siegel, Marvin
TITLE OF INVENTION: HEPATITIS B VIRUS TREATMENT
FILE REFERENCE: 12071-017002
CURRENT APPLICATION NUMBER: US/10/068,059
CURRENT FILING DATE: 2002-06-04
PRIOR APPLICATION NUMBER: US 60/266,733
PRIOR APPLICATION NUMBER: US 60/266,733
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8, Application US/10068059
Patent No. US20020155434A1
GENERAL INFORMATION:
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US-10-068-059-8
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Best Local (
                                                              Query Match 67.2%;
Best Local Similarity 95.5%;
Matches 147; Conservative
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                                                                                                                                                                                      LENGTH: 170
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           136 LEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVNAK 172
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Local Similarity 93.6%;
les 147; Conservation
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28 MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL 87
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Pred. No. 2.8e-68;
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Pred. No. 5.9e-69;
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Search completed: February 3, 2003, 09:35:29	Db 136 LEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVV 169	•			
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Perfect score:
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2: /cgn2_6/ptodata/1/laa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/laa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/laa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/laa/pCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/laa/backfiles1.pep:*
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US-07-739-642-2
US-07-739-142-2
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US-09-248-588-4
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US-08-968-747-18
US-08-968-747-21
US-08-968-747-2
US-08-968-747-2
US-08-105-483-217
US-08-709-209-217
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Sequence 3, Appli
Sequence 18, Appl
Sequence 21, Appli
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Sequence 217, Appli
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Sequence 2, Appli
Sequence 12, Appli
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STATE: MA
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Result

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RESULT 1 US-08-968-747-, Sequence 3, Sequence 1, Sequence 1, Sequence 2, Patent NO. 6 GENERAL INP APPLICANT TITLE OF NUMBER OF NUMBER OF CORRESPON CORRESPON STREET:		4. 10	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	
ULT 1 08-968-747-3 08-968-747-3 08-968-747-3 equence 3, Applicati atent No. 6060595 GENERAL INFORMATION: APPLICANT: Scagli TITLE OF INVENTION NUMBER OF SEQUENCE CORRESPONDENCE ADD ADDRESSEE: Fish STREET: 225 FR		157	166	175	194	607.5	671		767.5	767.5	767.5	772.5	772.5	772.5	821.5	821.5	٠	٠		
SULT 1 -08-968-747-3 Sequence 3, Application US/08968747 Patent No. 6060595 GENERAL INFORMATION: APPLICANT: Scaglioni et al. TITLE OF INVENTION: INHIBITION NUMBER OF SEQUENCES: 21 CORRESPONDENCE ADDRESS: ADDRESSEE: Fish & Richardson STREET: 225 Franklin Street		13.7	14.5	15.2	16.9	52.9	•	٠	•		66.9	67.3	67.3	67.3	71.6	71.6	71.6	73.6	75.2	
Sation US/0896874 SON: ION: et al. ION: INIBITION INCES: 21 ADDRESS: 1'sh & Richardson Franklin Street		420							214	214	214	214	214	214			4	159	351	
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US/08968747 et al. et al. INHIBITION OF VIRAL REPLICATION 21 S: S: Richardson P.C.	ALIGNMENTS	PCT-US96-10602-10	US-09-248-588-13	US-09-248-588-11	US-08-968-747-5	PCT-US96-10602-2	US-09-248-588-7	US-09-248-588-9	US-07-739-142-6	US-07-739-643-6	US-07-739-642-6	US-07-739-142-12	US-07-739-643-12	US-07-739-642-12	US-08-968-747-19	US-08-968-747-17	US-08-968-747-1	US-08-445-585-3	PCT-US96-10602-4	
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		Appl Appl	1001	11000	11001	11001	TIGGE	TOOM	70011	Appli	Tddy					100	10011	10011	Appli	

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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS;
LENGTH: 212 amino acids
TYPE: amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                        Query Match
Best Local Similarity
Matches 197; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IEM PC compatible

OPERATING SYSTEM: Windows

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/968,747

FILING DATE: 03-SEP-1997

CLASSIFICATIN: 435
                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEPAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 084
EHCSPHHTALRQAILCWGELWTLATWVGVNLEDPEFRGDASRDLVVSYVNTNWGLKFRQL 132
                                                                                PTVQASKLCLGMIMGMDIDPYKBFGATVELLSFLPSDFFPSVRDLLDTASALYREALBSP
                                                          PTVQASKLCLGWLWGWDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP 74
                                                                                                                                                                                                                                                                                       212 amino acids
                                                                                                                                          Conservative
                                                                                                                                                          90.6%;
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Pred. No. 2.2e-106;
0; Mismatches 1;
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                                                                                                                                                                       Length
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RESULT 2
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                                                                                                                                                                                                                                                                                                                          ; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-968-747-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/968,747
FILING DATE: 03-SEP-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 08472/705001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 192; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 617/542-50
TELEPAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Scaglioni et al. TITLE OF INVENTION: INHIBIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 194 amino acid
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                                                                                                                                62
                                                                                                                                                             78
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT 192
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                                                                                    SCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRTFSPRR 197
                                                                                                                             HHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKERQLLWFHI 137
                                                                                                                                                                                              SKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSP 61
                                                               SCLTEGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRTPSPRR 176
                                RRSQSPRRRRSQSREPQC 215
 RRSÓSPRRRRSÓSRESOC 194
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GENERAL INFORMATION:
APPLICANT: Scagli
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Hest Local Similarity 96.5%;
Matches 191; Conservation
                                                                                                                                                         Sequence 2, Application US/08968747 Patent No. 6060595
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SEQUENCE CHARACTERISTICS:
LENGTH: 199 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 0847
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5970
TELEFAX: 617/542-8906
TELEFAX: 60154
                                                                                                                                           GENERAL INFORMATION:
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TITLE OF INVENTION: INHIBITION OF VIRAL REPLICATION
NUMBER OF SEQUENCES: 21
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
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                                                                                        APPLICANT: Scaglioni et al.
TITLE OF INVENTION: INHIBITION OF VIRAL REPLICATION
NUMBER OF SEQUENCES: 21
                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C
AUL...
STREET: ...
CITY: Boston
Te: MA
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FILING DATE: 03-SEP-1997
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ZIP: 02110-2804
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                                                         ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                            HHTALRQAILCWGBIMTLATWVGVNLEDYKDDDDKSRDLVVSYVNTNWGLKFRQLLWFHI
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225 Franklin Street
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                                           225 Franklin Street
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Pred, No. 1.6e-103;
1; Mismatches 6;
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RESULT 5
US-08-105-483-217
; Sequence 217, Application US/08105483
neart No. 5494807
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US-08-968-747-2
                                                                                                                                                                                                                                                      Patent No. 5494807
Patent No. 5494807
GENERAL INFORMATION:
APPLICANT: Paoletti, Enzo
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN
TITLE OF INVENTION: 462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPONE: 617/5%.
TELEPAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
FRIGHT: 193 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 87.2%;
Best Local Similarity 96.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IMP PC compatible
COMPUTER: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/105,483
                                                                                                                                              STREET: 5..
STREET: 5..
CITY: New York
TATE: NY
                                                                                                                                                                                                          NUMBER OF SEQUENCES: 462
CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris & Safford
ADDRESSEE: C/o William S. Frommer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKST NUMBER: 084
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/968,747 FILLING DATE: 03-SEP-1997 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   116
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows
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                                                                                                                                            COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 193 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SCLTFGRETVIEYLVSEGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRTPSPRR 197
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                                                                                                                                                                                                 530 Fifth Avenue
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Pred. No. 4.7e-102;
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                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/709,209
FILING DATE: 21-AUG-1996
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/105,483
PRICATION NUMBER: US 08/105,483
PRICATION NUMBER: US 08/105,483
PRILING DATE: 12-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 217, Application US/08709209 Patent No. 5762938
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Best Local Similarity 88.7%;
Matches 189; Conservative
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APPLICANT: Paolet
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                                                                                                                                                                                                                                                                                        STREET: 550
CITY: New York
CTATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 346 amino acids
TYPE: amino acid
STRANDEDNESS: single
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APPLICATION NUMBER: US 07/847,951
FILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Prommer, William S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2400
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US 07/847,951 FILING DATE: 06-MAR-1992
                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Curtis, Morris & Safford ADDRESSEE: c/o William S. Frommer STREET: 530 Fifth Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TMMGLKFRQLLWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVR
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Pred. No. 1.5e-98;
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APPLICANT: No. 5766599ton, Elizabeth K.
APPLICANT: Riviere, Michel
APPLICANT: de Taisne, Charles
APPLICANT: de Taisne, Charles
APPLICANT: Limbach, Keith J.
APPLICANT: Donneon, Gerard P.
APPLICANT: Donneon, Gerard P.
APPLICANT: Cox, William I.
APPLICANT: Audonnet, Jean-Christophe Francis
APPLICANT: Rudsell Robert
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN
NUMBER OF SEQUENCES: 467
CORRESSEE: Curtis, Morris & Safford
ADDRESSEE: Curtis, Morris & Safford
ADDRESSEE: Cortis, Morris & Safford
ADDRESSEE: 530 Fifth Avenue
CITY: New York
Graffer. No
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 346 amino acids
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APPLICANT:
APPLICANT:
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REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2400
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
              ZIP: 10036

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/458,101

FILING DATE: 01-JUN-1995
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TYPE: amino acid
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Local Similarity 88.7%;
CLASSIFICATION:
                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enzo
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pred. No. 1.5e-98;
2; Mismatches 6
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REFERENCE/DOCKET NUMBER: 4543:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEPAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 217:
SEQUENCE CHARACTERISTICS:
LENGTH: 346 amino acids
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APPLICANT: RUTTER, WILLIAM J.; GOODMAN, HOWARD

APPLICANT COMMAND HEBAIT
                                                                                                                                                                                                                                                                                                              SEQ ID NO:13:
LENGTH: 211
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                                                                                                                                                                                                                                 Matches 187;
                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/07/679,621
PRILING DATE: 7-DEC-1984
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 513,055
FILING DATE: 12-JUL-1983
APPLICATION NUMBER: 107,267
FILING DATE: 21-DEC-1979
APPLICATION NUMBER: 41,909
FILING DATE: 24-MAY-1979
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: VACCII NUMBER OF SEQUENCES: 21 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     183 RRGRSPRRTPSPRRRRSQSPRRRRSQSRBPQC 215
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                              63 ALYRBALESPEHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVN 122
191 RTPSPRRRRSQSPRRRRSQSREPQC 215
                                                                                                                                                              3 LSSIFSRIGDPTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTAS
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                                                                                                                 EHCSPHHTALROAILCWGELMTLATWYGVNLEDPBERGDASRDLVVSYVNTNMGLKFRQL 132
                                                                                               EHCSPHHTALRQAILCWGEL-TLATWVGNNLEDP----ASRDLVVNVVNTNVGLKIRQL 126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VACCINES
                                                                                                                                                                                                                                                 83.1%; Score 954.5; DB 6;
91.2%; Pred. No. 6e-97;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 972; DB 1; Length 346; Pred. No. 1.5e-98; 2; Mismatches 6; Indels
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                                                                                                                                                                                                                                3; Mismatches
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; ORGANISM: Hepatitis B virus
US-09-248-588-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 12, Application PC/TUS9610602 GENERAL INFORMATION:
                                                                                                                      COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 182; Conserv
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LENGTH: 18
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TITLE OF INVENTION: Strategically Modified Hepatitis B Core Proteins and TITLE OF INVENTION: their Derivatives
FILE REFERENCE: SYN-101 4564/69529

CURRENT APPLICATION NUMBER: US/09/248,588

CURRENT FILING DATE: 1999-02-11

EARLIER APPLICATION NUMBER: 60/074537

EARLIER FILING DATE: 1998-02-12

SOFTWARE: Patentin Ver. 2.0
              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/017,814
FILING DATE: 20-JUN-1995
                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: The General Hospital Corporation
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           176
                                                                        APPLICATION NUMBER:
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                              CITY: Boston
STATE: MA
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CLASSIFICATION:
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                                                                                                        PCT/US96/10602
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Pred. No. 1.1e-96;
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                                                                                                                                            Version #1.30
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US-08-968-747-20
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     TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: ;
SEQUENCE CHARACTERISTICS:
LENGTH: 183 amino acids
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                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: Windows
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/968,747
FILING DATE: 03-SEP-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 08472/705001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEPAX: 617/542-5076
TELEPAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20, Application US/08968747 Patent No. 6060595
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' Tocal Similarity
Conserv
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APPLICANT: Scagilioni et al.
TITLE OF INVENTION: INHIBITION OF VIRAL REPLICATION
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 183 amino acids
TYPE: amino acid
TOROLOGY. 13001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 617/
TELEX: 200154
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amino acids
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95.7%;
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Pred. No. 2.8e-95;
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; STRANDEDNESS: not re.
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-968-747-20
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PCT-US96-10602-6
                                                                    MOLECULE TYPE: protein PCT-US96-10602-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 81.6
Best Local Similarity 95.7
Matches 180; Conservative
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Owery Match
Best Local Similarity
Matches 179; Conserv
                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 0078
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617,542-5070
                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: The General Hospital Corporation TITLE OF INVENTION: INHIBITION OF HEPATITIS NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: #loppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
COURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/10602
                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/017,814
FILING DATE: 20-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                176
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                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 397 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28 MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL 87
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: MA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 225 |
CITY: Boston
                                                                                                       TOPOLOGY:
                                                                                                                                                                                      TELEFAX: 617/
TELEX: 200154
                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                    ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CWGELWTLATWVGVNLEDP----ASRDLVVSYVDTNMGLKFRQLLWFMISCLTFGRETV 115
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                                                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 225 Franklin Street
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linear
                                                                                                                                                                                                       617/542-8906
          Conservative
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                                                                                                       linear
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95.7%;
          81.3%; Score 933.5; DB 5; 97.3%; Pred. No. 3e-94; tive 0; Mismatches 0;
                                                                                                                                                                          6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     B REPLICATION
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                                          Length 397;
               Indels
               <u>ა</u>
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US-07-739-642-2
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Patent No. 5173427
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APPLICANT: Mallonee, Richard L.
TITLE OF INVENTION: Vectors And Hosts With Increased
TITLE OF INVENTION: Expression Of HBcAg
NUMBER OF SEQUENCES: 24
                                                                                                                                                                                               Matches 178;
                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                             TELEFAX: 201-848-9228
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 07417-1880
ZIP: 07417-1880
COMPUTER READABLE FORM:
MEDIUM TYPE: P10ppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/739,642
FILING DATE: 1991881
                                                                                                                                                                                                                                                                                                                                                         NAME: Stierwalt, Brian K.
REGISTRATION NUMBER: 33,213
REFERENCE/DOCKET NUMBER: P-2272
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-848-5317
TELEFAX: 201-848-9228
                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 185 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      208
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116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Richard R. I
STREET: 1 Becton Drive
                                                                                                                                               28 MDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESPEHCSPHHTALRQAIL 87
                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                 TOPOLOGY:
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LEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRDRGRSPRRRTPSPRRRRSQSPRR
                                                                                 CWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFRQLLMFHISCLTFGRETV 147
                                                             CWGELMTLATWVGNNLEDP----ASRDLVVNYVNTNMGLKIRQLLWFHISCLTFGRETV 115
                                                                                                                              MDÍDÞYKEFGATVELLSFLÞSDFFÞSVRDLLDTASALYREALESÞEHCSPHHTALRQAIL 60
                 IEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRR--RGRSPRRRTPSPRRRRSOSPRR 205
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93.7%;
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Pred. No. 1.6e-93;
2; Mismatches 3
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206 RRSQSREPQC 215

US-07-739-643-2

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RESULT 15
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 185 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
Sequence 2, Application US/07739142
Patent NO. 5175272
GENERAL INFORMATION:
APPLICANT: Mallonee,, Richard L.
TITLE OF INVENTION: DNA Sequences With Increased Expression
TITLE OF INVENTION: of HECAG
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 80.4%; Score 922.5; DB 1; Length 185; Best Local Similarity 93.7%; Pred. No. 1.6e-93; Matches 178; Conservative 2; Mismatches 3; Indels 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/739,643
FILING DATE: 19910801
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Stierwalt, Brian K.
REGISTRATION NUMBER: 33,213
REFERENCE/DOCKET NUMBER: P-2090
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-849-5317
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TITLE OF INVENTION: Increased Expression of HBcAg
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
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TELEPHONE: ZUL-U-
TELEPHONE: ZUL-U-
201-848-9228
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STREET: 1 Becton Drive
CITY: Franklin Lakes
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Search completed: February Job time : 14.1098 secs

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Best Local Similarity
Matches 178; Conserv
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ZIP: 07417-1880

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/739,142
FILING DATE: 19910801
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Stierwalt Marian K
                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 201-848-9228 INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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176 RRSOSRESOC 185
                                                         116 LEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRDRGRSPRRRTPSPRRRRSQSPRR 175
                               206 RRSQSREPQC 215
                                                                             148 IEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRR--RGRSPRRTPGPRRRRRGQSPRR 205
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STREET: 1 B
CITY: Frank
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                                                                                                                                                                                                                                                                                                                                                                    TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Stierwalt,, Brian K.
REGISTRATION NUMBER: 33,213
REFERENCE/DOCKET NUMBER: P-:
                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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Franklin Lakes
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93.7%;
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Pred. No. 1.6e-93;
2; Mismatches 3;
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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                BLOSUM62
Gapop 10.0 , Gapext 0.5
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1148
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1982.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1983.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1983.DAT:*
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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HBV core antigen e	Hepatitis B virus	בינו			Henatitis B virus		Hepatitis B virus		HBV genotype B pre	is B virus	٤,	,	3 0 (itia P	3	atitis B virus	enotype	itis B	18 B	8 8	itie B	itie B	itis B	18 8	18 8	7	0 0	n (בי בי	, i	י נים נים	י מ	3. E	itis B	Hepatitis B virus

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WPI; 2000-514959/46. N-PSDB; AAA71735.	Hildt E, Hofschneider P;	(HILD/) HILDT E.	05-FEB-1999; 99DE-1004800.	04-FEB-2000; 2000WO-DE00363.	10-AUG-2000.	WD200046376-A2.	Synthetic.	Hepatitis b virus.	large surface protein; core antigen; gene therapy.	virus protein; cell permeability; cell-specific binding site: The.	Fusion protein; protein coat; virus-specific packaging signal and	HBV fusion protein comprising HBcAg and RGD.	08-JAN-2001 (first entry)	AAB10597;	RESULT 1 AAB10597 ID AAB10597 standard; Protein; 215 AA. XX

Particle for cell-specific gene delivery, useful in gene therapy,

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel particle (A), comprising a protein coat with a fusion protein (FP), and, inside the coat, a nucleic acid (I) including the sequence for a virus-specific packaging signal (psi) and a structural gene. FP contains a virus protein (VP), a peptide (P) that mediates cell permeability and a heterologous cell-specific binding site (RGD). The invention also describes (I) producing (A) in which FP (CONTAINS an LHBS (large surface protein of hepatitis B virus (HBV)) and RGD; (2) preparing (A) in which FP contains an HBV core antigen (HEcAg), (P) and RGD, (3) FP; (4) DNA encoding FP; and (5) expression vector containing the DNA of (d). The producte of the invention are used in gene therapy of cells and tissues, in vivo or ex vivo. This sequence represents a fusion protein which is described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         comprises nucleic of viral protein,
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                                                                                                                                                                                                              Vaccine; molecular scaffold; pilus; pilin; HBcAg; antigen; hepatitis B virus capsid protein; JUN; FOS; HIV gp140; messles virus N protein; bee venom phospholipase; Th type 2 T-helper; Th2; Sinbis virus E2 protein; amyloid beta; influenza M2 antigen; human immunodeficiency virus infection; viral hepatitis; messles; chicken pox; pneumonia; tuberculosis; syphilis; malaria; allergy; cancer; chronic disease; arthritis; colitis; diabetes;
                                                                                                                                                                                                                                                                                                                                           Hepatitis B virus core antigen variant (HBcAg)
                                                                                                                                                                                                                                                                                                                                                                           09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                      AAU80921 standard; Protein; 212
                                                                                                                                                                     Hepatitis B virus.
                                                                            02-MAY-2001; 2001WO-1B00741.
                                                                                                                                        WO200185208-A2
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(CYTO-) CYTOS BIOTECHNOLOGY AG (SEBB/) SEBBEL P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                      sclerosis
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                                               2000US-202341P
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permeability peptide
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Pred. No. 2.5e-109;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              that includes a fusion protein and cell-binding site -
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AAW50250 ID AAW5 XX

AAW50250 standard;

Protein; 212 AA

192

129

132 74

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complecular scaffold (molecular scaffold) which comprises a core core particle such as a bacterial pilus or pilin protein, a recombinant form conficient (HBCAR), and an organiser; and (b) an antigen or antigenic correction (HBCAR), and an organiser; and (b) an antigen or antigenic correction (HBCAR), and an organiser; and (b) an antigen or antigenic determinant include JUN, FOS, HIV gpl40, measies virus N components include JUN, FOS, HIV gpl40, measies virus N components include JUN, FOS, HIV gpl40, measies virus N components include JUN, FOS, HIV gpl40, measies virus N components include JUN, FOS, HIV gpl40, measies virus N components in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the subject in the 
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Best Local 9
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(BACH/)
(TISS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New composition, useful for vaccine production, comprises antigen of antigenic determinant and non-natural molecular scaffold comprising organizer and core particle such as bacterial pilus or pilin protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 227-228; 287pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-055561/07
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                                                                                                        LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT
                                                                                                                                                                                    EHCSPHHTALROAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVVTNMGLKFROL
                                                                                                                                                                                                                                              PTVQASKLCLGWLWGMDIDPYKBFGATVELLSFLPSDFFPSVRDLLDTASALYRBALESP
                          PSPRRRRSQSPRRRRSQSREPQC
                                                                              LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT
                                                                                                                                                                EHCSPHHTALRQAILCWGELMTLATWVGVNLEDP
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) BACHMANN M.
) TISSOT A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ש
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LECHENER F.
                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                   212 AA;
                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       es to a composition comprising: (a) a non-natural (molecular scaffold) which comprises a core
                                                                                                                                                                                                                                                                                                                                                       91.3%;
97.5%;
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                                                                                                                                                                                                                                                                                                                                     Score 1048.5; DB:
Pred. No. 3.7e-99;
0; Mismatches 0
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                                                                                                                                                                       ASRDLVVSYVNTNMGLKFRQL
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                                                                                                                                                                                                                                                                                                                    This polypeptide comprises the hepatitis B virus (HBV) 25 kDa (p25) CC Evidence is provided that HBV replication is inhibited in the CC presence of high levels of HBV precore or precore-related proteins. CC These proteins can be incorporated into HBV nucleocapsids along the process of the precore protein (see AAM50251), which is the usual component, and thereby render the nucleocapsids (cc expression of the precore proteins, or certain variants of them, cc expression of the precore proteins, or certain variants of them, cc expression of the precore proteins, or certain variants of them, cc expression of the precore proteins, or certain variants of them, cc inhibitory proteins include p25, p22 (see AAW50241), Met-p22 (see AAW50242), p18 (see AAW50236), Met-p18 (see AAW50238). The inhibitory proteins can be produced by recombinant methods using claimed expression vectors and host cells. They can be provided exogenously to the target cells for construct that directs overexpression of an inhibitory protein in Cc construct that directs overexpression of an inhibitory protein in
                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 197
                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15; Page 35; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA encoding proteins which can be incorporated with wild type nucleocapsid subunit(s) into a viral nucleocapsid - useful for inhibition of viral replication, especially hepatitis B virus
190
                           193
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                                                          130
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Viral replication; inhibitor; HBV; nucleocapsid; gene therapy; hepatocyte; liver; p25 protein.
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            PSPRRRRSQSPRRRRSQSREPQC 215
                                                                  LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNADILSTLPETTVVRRRGRSPRRRT 192
                                               LWFHISCLTFGRETVIEYLVSFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSPRRRT
                                                                                                           HCSPHHTALROAILCWGELMTLATWYGVNLEDP----ASRDLVVSYVNTNMGLKEROL
                                                                                                                             EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKERQL 132
                                                                                                                                                                     FTVQASKLCLGWLWGWDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP 74
                                                                                                                                                                                            PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP
                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                       212
                                                                                                                                                                                                                                      90.6%;
nilarity 97.0%;
Conservative
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Pred. No. 2.5e-98;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HBcAg) fusion proteins and polynucleotides encoding such proteins. Sequences of the invention are useful in methods of prophylactic or therapeutic vaccination or to manufacture medicaments for prophylactic or or therapeutic vaccination of the human or animal body against HBV, e.g. against viral hepatitis. They are also useful as a prophylactic vaccine against e.g. hepatitis c virus, influenza, polio, herpes, rables, acquired immune deficiency syndrome (AIDS) or foot-and-mouth disease. The present sequence is hepatitis B virus core antigenic protein (HBCAg).
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis B (HB) core antigen fusion proteins, useful as vaccines for the prophylactic or therapeutic treatment of humans or animals against e.g. HB virus, viral hepatitis, hepatitis C virus, influenza, or foot-and-mouth disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peptide
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                                                                                                                                                                                                                                                                                          3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention relates to hepatitis B virus (HBV) core antigen
LWEHISCLTEGRETVIEYLVSEGVWIRTEPAYRPENAPILSTLEETTVVRRRGRSERRRT 192
                                                                                                                                          EHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEPRGDASRDLVVSYVNTNMGLKFRQL 132
                                                                                                                                                                                                                                                     PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP
                                                                                                                                                                                                                        PTVQASKLCLGWLWGMDIDPYKBPGATVELLSPLPSDFFPSVRDLLDTASALYREALESP
                                                                                                                   EHCSPHHTALRQAILCWGELMTLATWVGVNLEDP---
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DB; AAD31509.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= Mature_HBc_protein
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                                                                                                                                                                                                                                                                                                                                                               90.6%; Score 1040.5; DB 97.0%; Pred. No. 2.5e-98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a composition comprising ribavirin and an antigen preferably non structural 3 protein (NS3)/4A fragment of hepatitis C virus (HCV) genome or a peptide or nucleic acid of HCV sequence. The composition is useful for enhancing an immune response to a hepatitis C antigen in humans, domestic, sport or pet species and as vaccines for treating and preventing HCV infections. The composition is also useful for treating viral, bacterial, fungal diseases and cancer. The present sequence is Hepatitis B virus C antigen (HBcAg) and E antige (HBeAg) sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-AUG-2000; 2000US-225767P
29-AUG-2000; 2000US-229175P
03-NOV-2000; 2000US-0705547
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vaccine compositions for treating and preventing disease, preferably hepatitis C virus infection, comprises ribavirin and antigen that has epitope present in hepatitis C virus -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 11; Page 82; 120pp; English.
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                                                                                             EHCSPHHTALROAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNMGLKFROL 132
                                                                                                                                                                                                                                                                                                     PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP 74
                                                                                                                                                                                                                                                                                                                                PTVQASKLCLGWLWGWDIDFYKEFGATVELLSELPSDFF5SVRDLLDTASALYREALESP 72
                                     PSPRRRRSQSPRRRRSQSREPQC 215
PSPRRRRSQSPRRRRSQSRESQC
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 1040.5;
Pred. No. 2.5
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                                                                                                                                                                                                                                                                                                                                                       carrier to present the epitope to the immune system. This chimeric protein or its nucleic acid is useful as a vaccine or in a method of protein or its nucleic acid is useful as a vaccine or in a method of prophylactic or therapeutic vaccination of the human or animal body, particularly against HBV. The nucleic acid encoding the protein may be used in gene therapy or DNA vaccination protocols. The chimeric protein or its nucleic acid may also be used as the basis of a prophylactic vaccine against a range of diseases, e.g. HBV, hepatitis prophylactic vaccine against a range of diseases, e.g. HBV, hepatitis A virus (HAV), hepatitis C virus (HCV), influenza, foot-and-mouth A virus (polio, herpes, rabies, acquired immunodeficiency syndrome (AIDS), dengue fever, yellow fever, malaria, tuberculosis, whooping cough, salmonellosis, typhoid, food poisoning, diarrhoea, meningitis or gonorrhea. The present sequence is Hepatitis B virus core antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New proteins comprising a modified hepatitis B core antigen, usefu
a vaccine in prophylactic or therapeutic vaccination of the human
animal body, particularly against hepatitis B virus infection -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-JUN-2000; 2000GB-0015308
06-OCT-2000; 2000GB-0024544
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to modified proteins comprising hepatitis B virus (HBV) core antigen (HBcAg) wherein one or more of the four arginine repeats has been deleted and the protein comprising the C-terminal cysteine of HBcAg. The deleted region may be replaced by an epitope from a protein other than HBcAg, in which case the HBcAg acts as a carrier to present the epitope to the immune system. This chimeric
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 39-40; 40pp; English.
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                                                                                                                                                                                                                                                                 Sequence
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PTVQASKLCLGWLWGWDIDPYKEFGATVELLSFLPSDFPFSVRDLLDTASALYREALESP 72
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                                                                                                                                                                                                                                                                     212 AA;
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Pred. No. 2.5
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RESULT 7

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XX VIN

PR 07-1

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XX Hepg

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CC Tinc

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                                                                                                                                                             The invention relates to the complete nucleic acid sequence of a new channan hepatitis B virus (HBV) genotype, provisionally named genotype CG. This genotype was found with a high prevalence in patients chronically infected with HBV and residing in Europe and the USA. The invention relates to a fully defined sequence of 3248 nucleotides as given in specification, a sequence with 924 identity to the given sequence, or sequence that is degenerate to the mentioned sequences. These polymucleotides are useful for detecting antibodies in encoded by the polymucleotides are useful for detecting antibodies in directed against the proteins are useful for detecting antibodies and for detecting HBCAG and HBCAG in precore precursor proteins. They have infections. The preparing a vaccine or medicament for treating sequence alignment of the precore and core regions of the different the genetypes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated and/or purified hepatitis B virus polypeptide and polynucleotide sequences that are phylogenetically different from genotype A-F molecules, useful for HBV diagnosis, prophylaxis and
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                                                                                                                                       New composition, useful for vaccine production, comprises antigen or antigenic determinant and non-natural molecular scaffold comprising organizer and core particle such as bacterial pilus or pilin protein
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                                                                                                                                                                                                                                       ) BACHMANN M.
) TISSOT A.
) LECHENER F.
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The invention relates to a composition comprising: (a) a non-natural molecular scaffold (molecular scaffold) which comprises a core particle such as a bacterial pilus or pilin protein, a recombinant for of the protein, a virus-like particle or a hepatitis B virus capsid protein (HBcAg), and an organiser; and (b) an antigen or antigenic determinant, where the molecular scaffold and antigenic determinant interact to form an ordered and repetitive antigen array. Suitable antigenic determinants include JUN, FOS, HIV 99140, measles virus N

torm

phospholipase, Sinbis virus

М 2

antigenic determinant and organizer and core particl

Disclosure; Page 222; 287pp; English.

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derived peptides and influenza M2 antigen. The composition (or vaccine) is useful for immunisation, by administration to a subject, where the composition produces an immune response, such as humoral, cellular or protective immune response, preferably a Th type 2 T-helper (Th2) cresponse that is specific for the antigenic determinant. The administration induces antibodies specific for the antigenic determinant of a subtype corresponding to the Th2 subtype in the subject of a subtype corresponding to the Th2 subtype in the subject of a subtype corresponding to the Th2 subtype in the subject of a subtype corresponding to the Th2 subtype in the subject of production of vaccines for prevention of infectious diseases such as production of vaccines for prevention of infectious diseases such as production of vaccines for prevention of infectious diseases such as composition and for treating allergy, cancer, and chronic diseases induced or accelerated by a Th1 type and specific immune response of the Th2 type and allows the creation of highly efficient vaccines against infectious diseases, and creation of highly efficient vaccines against infectious diseases, and creation of highly efficient vaccines against infectious diseases, and correct on the type immune response. The present sequence is a peptide or accelerated by a Th1 type immune response. The present sequence is a peptide or accelerated correction incorporated into the compositions of the invention.
RESULT 9
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cc molecular scaffold (molecular scaffold) which comprises a core comparticle such as a bacterial pilus or pilin protein, a recombinant form cc particle such as a bacterial pilus or pilin protein, a recombinant form cc of the protein (HBCAg), and an organiser; and (b) an antigen or antigenic cc protein (HBCAg), and an organiser; and (b) an antigen or antigenic cc interact to form an ordered and repetitive antigen array. Suitable cc interact to form an ordered and repetitive antigen array. Suitable cc interact to form an ordered and repetitive antigen array. Suitable cc determinant include JUN, FOS, HIV gpl40, measles virus N cc derived peptides and influenza M2 antigen. The composition (or vaccine) cc derived peptides and influenza M2 antigen. The composition (or vaccine) cc administration produces an immune response, such as humoral, cellular or cc administration induces antibodies specific for the antigenic determinant. The cc protective immune response, preferably a Th type 2 T-helper (Th2) composition of vaccines antibodies specific for the antigenic determinant composition of a subtype corresponding to the Th2 subtype in the subject. The subject does not generate a Th2 subtype that is specific for pilus or pilin cc does not generate a Th2 subtype that is specific for pilus or pilin cc production of vaccines for prevention of infectious diseases such as cc production of vaccines for prevention of infectious diseases such as cc production of vaccines sistem, viral hepatitis, measles, chicken pox, cc preunonis, tuberculosis, syphilis, malaria, and for treating allergy, cancer, and chronic diseases induced or accelerated by a Th1 type cancer, and chronic diseases induced or accelerated by a Th1 type and allows the composition is useful to generate defined self-specific commune response. The present sequence is a peptide or protein incorporated into the compositions of the invention.
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                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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(SEBB/)
(DUNA/)
(BACH/)
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                                                                                                                                                                                                                                                                                                                                                                Sequence
190
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                                                                                           LWFHISCLTFGRETVIEYLVGFGVWIRTPPAYRPPNAPILSTLPETTVVRRRGRSFRRRT 192
                                                                                                                                                                                                                 PTVQASKLCLGWLWGWDIDPYKBFGATVELLSFLPSDFFPSVRDLLDTASALYREALESP 74
                                                                                                                                                                                                                                          PTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFPPSVRDLLDTASALYREALESP 72
                                                                                                                                                                EHCSPHHTALRQAILCWGELMTLATWYGYNLEDPEFRGDASRDLVVSYVNTNWGLKFRQL 132
                                                                                                                                              PSPRRRRSQSPRRRRSQSRESQC 212
                      PSPRRRRSQSPRRRRSQSRBPQC 215
                                                                    SEBBEL P.
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                                                                                                                                                                                                                                                                                                                                                                    212 AA;
                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                              90.4%;
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                                                                                                                                                                                                                                                                                           Score 1037.5; DE pred. No. 5e-98; 1; Mismatches
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AAU80916

standard;

Protein; 212

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CC The invention relates to a composition comprising: (a) a non-natural CC particle such as a bacterial pilus or pilin protein, a recombinant form CC of the protein, a virus-like particle or a hapatitis B virus capsid CC of the protein, a virus-like particle or a hapatitis B virus capsid CC of the protein, where the molecular scaffold and antigen or antigen continued to form an ordered and repetitive antigen array. Suitable CC interact to form an ordered and repetitive antigen array Suitable CC derived peptides and influenza M2 antigen array suitable CC is useful for immunisation, by administration to a subject, where the CC is useful for immune response, such as humoral, cellular or cresponse that is specific for the antigenic determinant. The composition of a subtype corresponding to the Th2 subtype in the subject. The subject codes not generate a Th2 subtype that is specific for the antigenic determinant CC of a subtype corresponding to the Th2 subtype in the subject. The subject corresponder of vaccines for prevention of infectious diseases such as humoral, cellular or CC response that is specific for the antigenic determinant. The composition is useful for the CC polypeptide or antigenic determinant. The composition is useful for the CC polypeptide or antispenic determinant. The composition is useful for the CC under, and chronic diseases induced or accelerated by a Th1 type and allows the CC cancer, and chronic diseases induced or accelerated by a Th1 type and allows the CC creation of highly efficient vaccines of the Th2 type and allows the CC by a Th1 type immune response. The present sequence is a peptide or accelerated corrected by a Th1 type immune response. The present sequence is a peptide or accelerated corrected by a Th1 type immune response. The present sequence is a peptide or accelerated corrected by a Th1 type immune response.
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organizer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              composition, useful for vaccine production, igenic determinant and non-natural molecular anizer and core particle such as bacterial pi
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scaffold comprising
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RESULT 11
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Best Local
                                          New composition, useful for vaccine production, comprises antigen or antigenic determinant and non-natural molecular scaffold comprising organizer and core particle such as bacterial pilus or pilin protein -s
                  Disclosure; Page 226;
                                                                                                                                                                                                                                                                                                                                     Vaccine; molecular scaffold; pilus; pilin; HBcAg; antigen; hepatitis B virus capsid protein; JUN; FOS; HIV gp140; measles virus N protein; bee venom phospholipase; Th type 2 T-helper; Th2; Sinbis virus E2 protein; amyloid beta; influenza M2 antigen; human immunodeficiency virus infection; viral hepatitis; measles; chicken pox; pneumonia; tuberculosis; syphilis; malaria; allergy; cancer; chronic disease; arthritis; colitis; diabetes;
                                                                                                                   Sebbel P,
                                                                                                                                                                           (SEBB/)
                                                                                                                                                                                                                    05-MAY-2000; 2000US-202341P
                                                                                                                                                                                                                                       02-MAY-2001; 2001WO-IB00741
                                                                                                                                                                                                                                                                                                           Hepatitis B virus.
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                                                                                                                                                                (BACH/)
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                                                                                              2002-055561/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PTVQASKLCLGWLWGMDIDPYKBFGATVELLSFLPSDFFPSVRDLLDTASALYRBALESP 72
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                                                                                                                                         LECHENER F.
                                                                                                                                                    BACHMANN M.
TISSOT A.
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                 287pp; English.
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                                                                                                                 Bachmann M,
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Pred. No. 5e-98;
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The invention relates

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composition comprising:

a)

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AAU80920
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                                                                        chicken pox; pneumonia; tuberculosis; syphilis; malaria; allergy; cancer; chronic disease; arthritis; colitis; diabetes;
                                                                                                                                                                                                                                       Hepatitis B virus core antigen variant (HBcAg) #25
                                                                                                                                                                                                                                                                                                                                                               AAU80920 standard; Protein;
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                Hepatitis B
                                                           multiple
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Pred. No. 8e-98;
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determinant, where the molecular scaffold and antigenic determinant, where the molecular scaffold and repetitive antigen array. Suitable contributed to form an ordered and repetitive antigen array, Suitable contributed to form an ordered and repetitive antigen array. Suitable contributed to form an ordered and repetitive antigen array. Suitable contributed between phospholipase, Sinbis virus E2 protein, amyloid beta corrective immunisation, by administration to a subject, where the contributed peptides and influenza M2 antigen. The composition (or vaccine) contributed antiperior to a subject, where the contributed immune response, preferably a Th type 2 T-helper (Th2) contributed that is specific for the antigenic determinant. The composition is useful for the subject of a subtype corresponding to the Th2 subtype in the subject. The subject contributed are a Th2 subtype that is specific for pillus or pi
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) DUNANT N.
) BACHMANN M.
) TISSOT A.
                                                                                EHCSPHHTALROAILCWGELMTLATWVGVNLEDP----
                                                                                                      EHCSPHHTALROAILCWGELMTLATWVGVNLEDPEFRGDASRDLVVSYVNTNWGLKFROL
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cc molecular scaffold (molecular scaffold) which comprises a correct particle such as a bacterial pilus or pilin protein, a recombinant form cc of the protein, a virus-like particle or a hepatitis B virus capsid correct protein, a virus-like particle or a hepatitis B virus capsid correct protein, a virus-like particle or a hepatitis B virus capsid correct protein, and an organiser; and (b) an antigen or antigenic determinant where the molecular scaffold and antigenic determinant capsid correct to form an ordered and repetitive antigen array. Suitable protein, bee venom phospholipase, Sinbis virus R2 protein, amyloid beta derived peptides and influenza M2 antigen. The composition (or vaccine) administration produces an immune response, such as bubject, where the protective immune response professionse, such as bumoral, cellular or response that is specific for the antigenic determinant. The administration induces antibodies specific for the antigenic determinant of a subtype corresponding to the Th2 subtype in the subject. The subject polypeptide or antigenic determinant. The composition is useful for the production of vaccines for prevention of infectious diseases such as human immunodeficiency virus, viral hepatitis, measles, chicken pox.
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                                                                                                                                                                                                                                                                                                                                The invention relates to a composition comprising: (a) a non-natural molecular scaffold (molecular scaffold) which comprises a core
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DUNANT N.
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TISSOT A.
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Matches 196;
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            WPI; 2002-055561/07
                                       Sebbel P,
                                                                                           (DUNA/)
                                                                                                                                                         05-MAY-2000; 2000US-202341P
                                                                                                                                                                                02-MAY-2001; 2001WO-IB00741.
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                                                                                                                                                                                                                                                                  Hepatitis B virus.
                                                                                                                                                                                                                                                                                          multiple sclerosis.
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cancer; chronic disease; arthritis; colitis; diabetes;
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                                                                                                       SEBBEL P.
DUNANT N.
                                                                LECHENER F
                                                                            BACHMANN M.
TISSOT A.
                                                                                                                                CYTOS BIOTECHNOLOGY
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                                      Dunant N,
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96.6%;
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determinant, where the molecular scaffold and antigenic determinant of interact to form an ordered and repetitive antigen array. Suitable protein, bee varom phospholipase, Sinbis virus E2 protein, anyloid beta composition produces an immune response, such as subject, where the cadministration produces an immune response, such as humoral, cellular or protective immune response, preferably a Th type 2 T-helper (Th2) cresponse that is specific for the antigenic determinant. The camposition induces antibodies specific for the antigenic determinant. The composition is useful for a subtype corresponding to the Th2 subtype in the subject. The subject of a subtype corresponding to the Th2 subtype in the subject. The subject of a subtype corresponding to the Th2 subtype in the subject of production of vaccines for prevention of infectious diseases such as for prevention of infectious diseases such as the composition is useful for the cancer, and chronic diseases induced or accelerated by a Th1 type cancer, and specific immune response, such as arthritis, colitis, diabetes and multiple cantibodies and specific immune response of the Th2 type and allows the creating allergy, cancer, and chronic diseases induced or accelerated by a Th1 type immune response. The present sequence is a peptide or protein incorporated into the compositions of the inventious diseases.
  AAU80924
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Vaccine; molecular scaffold; pilus; pilin; HBcAg; antigen; hepatitis B virus capsid protein; JUN; FOS; HIV gpl40; measles virus N protein; bee venom phospholipase; Th type
                                                                                                                                                                                                         AAU80924 standard;
                                                                                                                                                                                                                                                                                                                                                 193
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                                                                                                                                                                                                                                                                                                                                                                                       EHCSPHHTALROAILCWGELWTLATWYGVNLEDPEFRGDASRDLVVSYVNTNMGLKFROL 132
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Query Match Best Local

Similarity

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90.1%; nilarity 96.1%; Conservative

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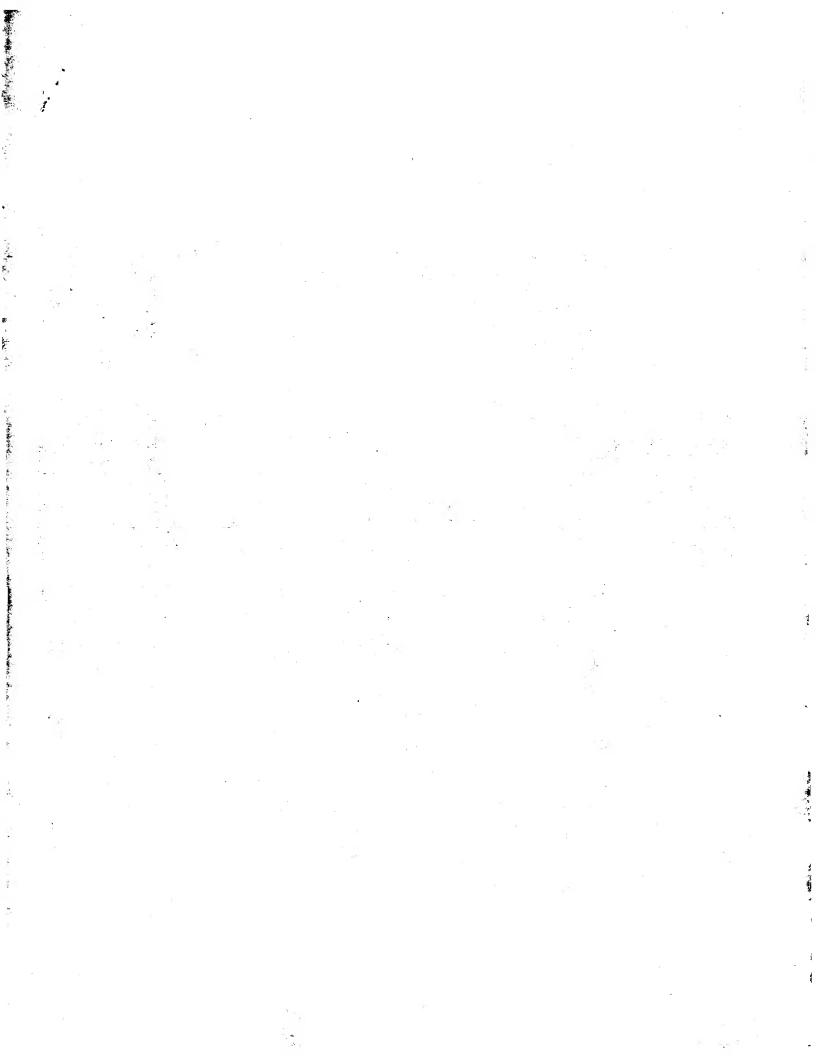
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CC determinant, where the molecular scaffold and antigenic determinant CC determinant, where the molecular scaffold and antigenic determinant (CC interact to form an ordered and repetitive antigen array. Suitable CC antigenic determinants include JUN, FOS, HIV gp140, measles virus N CC protein, bee venom phospholipase, Sinbis virus E2 protein, amyloid beta CC derived peptides and influenza M2 antigen. The composition (or vaccine) CC is useful for immune response M2 antigen. The composition (or vaccine) CC response that is specific for the antigenic determinant. The composition induces antibodies specific for the antigenic determinant. The CC des not generate a Th2 subtype that is specific for the antigenic determinant administration of vaccines subtype that is specific for pilus or pilus or polypetide or antigenic determinant. The composition is useful for the subject of production of vaccines for prevention of infectious diseases such as CC puspeptide or antigenic determinant. The composition is useful for the cancer, and chronic diseases induced or accelerated by a Th1 type CC cancer, and chronic diseases induced or accelerated by a Th1 type cancer, such as arthritis, colitis, diabetes and multiple collerated and specific immune responses of the Th2 type and allows the CC ereation of highly efficient vaccines against infectious diseases, and CC creation of highly efficient vaccines against infectious diseases, and CC protein incorporated into the compositions of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Th2; Sinbis virus B2 protein; amyloid beta; influenza M2 antigen; human immunodeficiency virus infection; viral hepatitis; measles; chicken pox; pneumonia; tuberculosis; syphilis; malaria; allergy; cancer; chronic disease; arthritis; colitis; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis B virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New composition, useful for vaccine production, comprises antigen or antigenic determinant and non-natural molecular scaffold comprising organizer and core particle such as bacterial pilus or pilin protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                particle such as a bacterial pilus or pilin protein, a recombinant for the protein, a virus-like particle or a hepatitis B virus capsid protein (HBcAg), and an organiser; and (b) an antigen or antigenic determinant, where the molecular scaffold and antigenic determinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a composition comprising: (a) a non-natural molecular scaffold (molecular scaffold) which comprises a core
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EMBL; X02496; CAA26324.1; ALT INIT.
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PIR; A03703; SAVLAH.
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Dishler A.V., Kalic Y.V., Pudova N.V., Gren
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MEDLINE=85204397; PubMed=3996597;
Bichko V., Pushko P., Dreilina D., Pumpen P., Gren E.Y.;
"Subtype ayw variant of hepatitis B virus. DNA primary structure analysis.";
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MEDLINE=81012091; PubMed=399327;

Galibert F., Mandart E., Fitoussi F.,

"Nucleotide sequence of the hepatitis
cloned in E. coli.";

Nature 281:646-650(1979).

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Major surface antigen precursor.
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MEDLINE=90266476; PubMed=2345966;

Tong S., Li J., Virviteki L., Trepo C.;

Tong S., Li J., Virus replication in the presence of anti-HBe

"Active hepatitis B virus replication in the presence of anti-HBe

associated with viral variants containing an inactive pre-C region
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Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
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Pasek M., Gotto T., Gilbert W., Zink B., Schall
Leadbetter G., Murray K.;
"Hepacitis B virus genes and their expression
Nature 282:575-579(1979).
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MEDLINE=85107103; PubMed=3968537;
Takeshima H., Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M., Shimada J., Omura Takeshima H. Inokoshi J., Namiki M. Inoko
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149 PISSIFSRIGDP 160
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Pred. No. 0.0012;
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cloned hepatitis B
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P12934; Q67860;
01-OCT-1989 (Rel. 12, Created)
F 01-UT-1989 (Rel. 12, Last sequence 15-JUL-1998 (Rel. 36, Last annotati
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Best Local Similarity
Matches 10; Conser
                                            CARBOHYD
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CARBOHYD
SEQUENCE
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or send a
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                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10409;
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Viruses; Retroid viruses; Hepadna
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PIR; A03705; SAVLA.
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X01587; CAA25743.1; ALT_INIT.
Pro; IPR000349; Hepvir_surfAg.
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      Similarity
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Pred. No. 0.37
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Pred. No. 0.01
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15-JUL-1998
15-JUN-2002
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SEQUENCE
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CARBOHYD
                                           Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                           Beta-defensin
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        SEQUENCE FROM N.A.
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                                                                                                                                                                                           161 ISSISSRIGDP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Domingo E., Mayumi M., "Nucleotide sequence of a hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis B virus (subtype adw / strain Philippino/pFDW294)
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus
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01-JUL-1993
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(Rel. 36, Last sequence update)
(Rel. 41, Last annotation update)
in 2 precursor (hBD-2) (Skin-antimicrobial peptide
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Pred. No.
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                                          Craniata; Vertebrata;
Catarrhini; Hominidae
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MEDLINE=97345625; PubMed=9202117;

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Harder J., "A peptide
                                                   EMBL; Z71389; CAA95992.1; -.
EMBL; AF040153; AAC33549.1; -.
EMBL; AF071216; AAC69554.1; -.
EMBL; AJ000152; CAB65126.1; -.
PDB; 1F03; 01-NOV-00.
PDB; 1F04; 01-NOV-01.
PDB; 1E4Q; 26-NOV-01.
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[2]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=99051334; PubMed=9831658;
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                                                                                                                                                                                                                                                                                                                                                                                     Kluever E., Schulz A., Foresmann W.-G. "Chemical synthesis of beta-defensins J. Pept. Res. 59:241-248(2002).
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                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for comentities requires a license agreement (See http://www.isb-sib.ch/anor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "The structure of human beta-defensin-2 order oligomerization.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    X-RAY CRYSTALLOGRAPHY (1.35 ANGSTROMS).
MEDLINE=20490730; PubMed=10906336;
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                                                                                                                                                                                                                                                                                                    -!- SUBCELLULAR LOCATION: Secreted
-!- TISSUE SPECIFICITY: EXPRESSED
-!- INDUCTION: BY INFLAMMATION.
                                                                                                                                                                                                                                                                                                                                            Protein Sci. 10:2470-2479 (2001).
                                                                                                                                                                                                                                                                                                                                                                   structural conservation in the absence similarity.";
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                                         HGNC: 2767;
IPR001855; Defensin_beta.
IPR001271; Defensin_mammal.
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and LEAP-1/hepcidin.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus isolated from a naturally infected J. Gen. Virol. 69:1383-1389(1988).
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Harrison T
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P17399;
01-AUG-1990
                                                                                                                                          "Typing hepatitis B virus by homology in comparison of surface antigen subtypes.";
J. Gen. Virol. 69:2575-2583(1988).
                        This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Enthe European Bioinformatics Institute. There are no rest
                                                                                                                                                                                                                              MEDLINE=89010694; PubMed=3171552;
Okamoto H., Tsuda F., Sakugawa H.,
Miyakawa Y., Mayumi M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CHAIN
CARBOHYD
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SEQUENCE
                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis B virus (subtype adw / strain Okinawa/pODW282).
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the EMPS person its supposed Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P17398;
01-AUG-1990
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PIR; G28925; SAVLJI. -
InterPro; IPR000349; Hepvir surfag.
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"Typing hepatitis B virus by homology in "Typing hepatitis B virus by homology in "Typing hepatitis B virus by homology in "Typing hepatitis B virus by homology in "Typing hepatitis B virus by homology in "Typing hepatitis" hepatitis B virus by homology in "Typing hepatitis" hepatitis 
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F., Sakugawa H.,
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Hepadnaviridae; Orthohepadnavirus
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RA Wood V., Gwilliam R., Rajandream M.A., Lyne M., Lyne R., Stewart A.,
RA Sgouros J., Peat N., Hayles J., Basham D., Bowman S.,
RA Brooks K., Brown D., Brown S., Chillingworth T., Churcher C.M.,
RA Brooks K., Brown D., Brown S., Chillingworth T., Churcher C.M.,
RA Gentles S., Goble A., Hamlin N., Harris D., Hidalgo J., Hodgson G.,
RA Holroyd S., Hornsby T., Howarth S., Huckle B.J., Hunt S., Jagels K.,
RA Holroyd S., Hornsby T., Howarth S., Huckle B.J., Hunt S., Jagels K.,
RA Holroyd S., Hornsby T., Howarth S., Huckle B.J., Hunt S., Jagels K.,
RA James K., Jones M., Leather S., McDean J.,
RA Oliver K., O'Neil S., Pearson D., Quail M.A., Rabbinowitsch E.,
RA Oliver K., O'Neil S., Pearson D., Quail M.A., Rabbinowitsch E.,
RA RICHERFORD K., Rutter S., Saunders D., Seeger K., Sharp S.,
RA Skilton J., Simmonds M., Squares R., Squares S., Stevens K.,
RA Taylor K., Taylor R.G., Tivey A., Walsh S.V., Warren T., Whitehead S.,
RA Woodward J., Volkaert G., Aert R., Robben J., Grymonprez B.,
RA Gabel C., Fuchs M., Fritzc C., Holzer E., Moestl D., Hilbert H.,
RA Gabel C., Fuchs M., Fritzc C., Holzer E., Moestl D., Hilbert H.,
RA Galbert F., Aves S. J., Xiang Z., Hunt C., Moore K., Hurst S.M.,
RA Berzym K., Langer I., Beck A., Lehracch H., Reinhardt R., Pohl T.M.,
RA Galbert F., Aves S. J., Xiang Z., Hunt C., Moore K., Hurst S.M.,
RA Lucas M., Rochet M., Gaillardin C., Tallada V.A., Garzon A., Thode G.,
RA Deminguez A., Revuelta J.L., Moreno S., Ammstrong J., Forsburg S.L.,
RA Cerrutti L., Lowe T., McCombie W.R., Paulsen I., Potashkin J.,
RT "The genome sequence of Schizosaccharomyces pombe.";
RL Nature 415:871-880(2002)
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Wood V., Gwilliam R., Rajandream N
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CARBOHYD
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erol O-acyltransferase (EC 2.3.1.26) (Sterol-ester
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scomycota; Schizosaccharomycetes;
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(By

LOCATION: Integral membrane protein. Endoplasmic By similarity).

BELONGS TO THE STEROL O-ACYLTRANSFERASE FAMILY.

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                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
STRAIND/UW-3/Cx;
MEDLINE=99000809; PubMed=9784136;
Stephens R.S., Kalman S., Lammel C.J., Fan J., Marathe R., Arayethens R.S., Kalman S., Lammel C.J., Fan Q., Koonin E.V.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Excinuclesse ABC subunit A.
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This SWISS-PROT entry is copyright. It is production the Swiss Institute of Bioinformatics
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Bacteria; Chlamydiales; Chlamydiaceae; Chlamydia.
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                                                                                                                                                                                                                                                                                                                               Chlamydia trachomatis."
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                                                                                                                                                   FUNCTION: THE ABC EXCISION NUCLEASE IS A DNA REPAIR ENZYME THAT CATALYZES THE EXCISION REACTION OF UV-DAMAGED NUCLEOTIDE SEGMENTS PRODUCING OLIGOMERS HAVING THE MODIFIED BASE(S). UVRA IS AN ATPASE AND A DNA-BINDING PROTEIN THAT PREFERENTIALLY BINDS SINGLE-STRANDED OR UV-IRRADIATED DOUBLE-STRANDED DNA (BY SIMILARITY). SUBUNIT: CONSISTS OF THREE SUBUNITS; UVRA, UVRB AND UVRC.
                                                                                                        SUBCELLULAR LOCATION: Cytoplasmic (By similarity).
SIMILARITY: BELONGS TO THE ABC TRANSPORTER FAMILY.
                                                                                    CONTAINS FOUR ABC DOMAINS.
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ProDom; PD000006; ABC_transportr;
TIGRPAM8; TIGR00630; uvra; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Caenorhabditis elegans.
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea;
Rhabditidae; Peloderinae; Caenorhabditis.
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PROSITE; PS00211; ABC_TRANSPORTER; FALSE_NEG.
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                                                                                                                                                                                                                                                                                                             or send an email to license@isb-sib.ch).
                                                                                                                                                                                           Hypothetical SEQUENCE 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (JAN-1995) to the EMBL/GenBank/DDBJ databases.
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5; Conserv
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495 AA; 55814 MW;
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(45.1 kDa protein F19C6.2 in chromosome
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Pred. No. 57;
2; Mismatches
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RESULT 15
DCUP CHAME
ID DCUP CHAME
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MVIN_SYNY3
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Best Local
                                                         DCUP_CHLMU
Q9PLH7;
16-OCT-2001
16-OCT-2001
15-JUN-2002
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
Uroporphyrinogen decarboxylase (EC 4.1.1.37) (URCHEME OR TC0123.
Chlamydia muridarum.
Bacteria; Chlamydiales; Chlamydiaceae; Chlamydia.
NCBI_TaxID=83560;
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                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR004268; MVIN like. Pfam; PF03023; MVIN; 1.
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30-MAY-2000
16-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SYMECHOCYBEIS BD. SETAIN PCC6803. I. Sequence features in the 1 region from map positions 64% to 92% of the genome."; DNA Res. 2:153-166(1995).

-i- SUBCELLULAR LOCATION: Integral membrane protein (Potential).

-i- SIMILARITY: BELONGS TO THE MVIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sugiura M., Tabata S.;
"Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PCC6803. I. Sequence features in the 1 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
MEDLINE=96127529; PubMed=8590279;
KANNEKO T., Tanaka A., Sato S., Kotani H.,
Sugiura M., Tabata S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synechocystis sp. (strain PCC 6803).
Bacteria; Cyanobacteria; Chroococcales; Synechocystis.
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Matches 6
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                                                                                                                                                                        PROSITE; PS00906; URÓD 1; 1.
PROSITE; PS00907; UROD 2; 1.
LYABE; Decarboxylase; Porphyrin biosynthesis; Complete proteome.
SEQUENCE 334 AA; 37380 MW; 199B6A88AC1F4D2D CRC64;
                                                                                                                                                                                                                                                TIGR; TC0123; -.
InterPro; IPR000257; Uro_decarbxyls.
Pfam; PF01208; URO-D; 1.
                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             -1- PATHWAY: Porphyrin biosynthesis.
-1- SUBCELLULAR LOCATION: Cytoplasmic (probable).
-1- SIMILARITY: BELONGS TO THE UROPORPHYRINGGEN DECARBOXYLASE FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic Acids Res. 28:1:
-I- CATALYTIC ACTIVITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Read T.D., Brunham R.C., Shen C., Gill S.R., Heidelberg J.F., White O., Hickey E.K., Peterson J., Utterback T., Berry K., Bass S., Linher K., Weidman J., Khouri H., Craven B., Bowman C., Dodson R., Gwinn M., Nelson W., DeBoy R., Kolonay J., McClarty G., Salzberg S.L., Eisen J., Fraser C.M.;
"Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.";
                                                             258 LSEIYRQLGDP
                                                                                                                        Local Similarity tes 6; Conserv
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                                                                                          LSSIFSRIGDP 12
                                                                                                                          Conservative
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Pred. No.
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